

Rong Zhong

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.

90
papers

1,525
citations

24
h-index

34
g-index

94
ext. papers

1,904
ext. citations

6.7
avg, IF

4.2
L-index

#	Paper	IF	Citations
90	Genome-wide gene-bisphenol A, F and triclosan interaction analyses on urinary oxidative stress markers. <i>Science of the Total Environment</i> , 2022 , 807, 150753	10.2	0
89	Urinary bisphenol A and its interaction with CYP17A1 rs743572 are associated with breast cancer risk. <i>Chemosphere</i> , 2022 , 286, 131880	8.4	1
88	No Evidence for a Causal Link between Serum Uric Acid and Nonalcoholic Fatty Liver Disease from the Dongfeng-Tongji Cohort Study.. <i>Oxidative Medicine and Cellular Longevity</i> , 2022 , 2022, 6687626	6.7	0
87	Hepatocellular carcinoma risk variant modulates lncRNA HLA-DQB1-AS1 expression via a long-range enhancer-promoter interaction. <i>Carcinogenesis</i> , 2021 , 42, 1347-1356	4.6	2
86	FOXA1 of regulatory variant associated with risk of breast cancer through allele-specific enhancer in the Chinese population. <i>Breast Cancer</i> , 2021 , 1	3.4	
85	rs641738 Is Not Associated With the Risk of Hepatocellular Carcinoma or Persistent Hepatitis B Infection. <i>Frontiers in Oncology</i> , 2021 , 11, 639438	5.3	1
84	Aberrant MCM10 SUMOylation induces genomic instability mediated by a genetic variant associated with survival of esophageal squamous cell carcinoma. <i>Clinical and Translational Medicine</i> , 2021 , 11, e485	5.7	2
83	A genetic variant conferred high expression of CAV2 promotes pancreatic cancer progression and associates with poor prognosis. <i>European Journal of Cancer</i> , 2021 , 151, 94-105	7.5	2
82	CancerImmunityQTL: a database to systematically evaluate the impact of genetic variants on immune infiltration in human cancer. <i>Nucleic Acids Research</i> , 2021 , 49, D1065-D1073	20.1	6
81	Identification of genetic variants in mA modification genes associated with pancreatic cancer risk in the Chinese population. <i>Archives of Toxicology</i> , 2021 , 95, 1117-1128	5.8	8
80	Colorectal cancer risk variant rs7017386 modulates two oncogenic lncRNAs expression via ATF1-mediated long-range chromatin loop. <i>Cancer Letters</i> , 2021 , 518, 140-151	9.9	1
79	Serum concentrations of organochlorine pesticides, biomarkers of oxidative stress, and risk of breast cancer. <i>Environmental Pollution</i> , 2021 , 286, 117386	9.3	5
78	Bisphenol A exposure, interaction with genetic variants and colorectal cancer via mediating oxidative stress biomarkers. <i>Environmental Pollution</i> , 2021 , 287, 117630	9.3	2
77	Associations of polycyclic aromatic hydrocarbons exposure and its interaction with XRCC1 genetic polymorphism with lung cancer: A case-control study. <i>Environmental Pollution</i> , 2021 , 290, 118077	9.3	1
76	Functional characterization of a low-frequency V1937I variant in FASN associated with susceptibility to esophageal squamous cell carcinoma. <i>Archives of Toxicology</i> , 2020 , 94, 2039-2046	5.8	5
75	Nonreceptor protein tyrosine phosphatases (NRPTs) gene family associates with the risk of hepatocellular carcinoma in a Chinese hepatitis B virus-related subjects. <i>Molecular Carcinogenesis</i> , 2020 , 59, 980-988	5	0
74	CpG-methylation-based risk score predicts progression in colorectal cancer. <i>Epigenomics</i> , 2020 , 12, 605-614	4.5	5

73	Risk SNP-Mediated Enhancer-Promoter Interaction Drives Colorectal Cancer through Both and. <i>Cancer Research</i> , 2020 , 80, 1804-1818	10.1	21
72	Genetic variants in m6A modification genes are associated with esophageal squamous-cell carcinoma in the Chinese population. <i>Carcinogenesis</i> , 2020 , 41, 761-768	4.6	21
71	A functional variant in TNXB promoter associates with the risk of esophageal squamous-cell carcinoma. <i>Molecular Carcinogenesis</i> , 2020 , 59, 439-446	5	2
70	Urinary bisphenol A and its interaction with ESR1 genetic polymorphism associated with non-small cell lung cancer: findings from a case-control study in Chinese population. <i>Chemosphere</i> , 2020 , 254, 126835	8.4	13
69	Genetic Predisposition to Colon and Rectal Adenocarcinoma Is Mediated by a Super-enhancer Polymorphism Coactivating and. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 850-859	4	4
68	LINC01149 variant modulates MICA expression that facilitates hepatitis B virus spontaneous recovery but increases hepatocellular carcinoma risk. <i>Oncogene</i> , 2020 , 39, 1944-1956	9.2	6
67	Evaluation of polymorphisms in microRNA-binding sites and pancreatic cancer risk in Chinese population. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 2252-2259	5.6	3
66	Regulatory Variant as Predictor of Epirubicin-Based Neoadjuvant Chemotherapy in Luminal A Breast Cancer. <i>Frontiers in Oncology</i> , 2020 , 10, 571517	5.3	2
65	ANKLE1 N -Methyladenosine-related variant is associated with colorectal cancer risk by maintaining the genomic stability. <i>International Journal of Cancer</i> , 2020 , 146, 3281-3293	7.5	15
64	Three functional variants were identified to affect RPS24 expression and significantly associated with risk of colorectal cancer. <i>Archives of Toxicology</i> , 2020 , 94, 295-303	5.8	5
63	Trans-acting non-synonymous variant of FOXA1 predisposes to hepatocellular carcinoma through modulating FOXA1-ER α transcriptional program and may have undergone natural selection. <i>Carcinogenesis</i> , 2020 , 41, 146-158	4.6	2
62	N-methyladenosine mRNA methylation of regulates AKT signalling to promote PTEN-deficient pancreatic cancer progression. <i>Gut</i> , 2020 , 69, 2180-2192	19.2	24
61	A functional variant in the boundary of a topological association domain is associated with pancreatic cancer risk. <i>Molecular Carcinogenesis</i> , 2019 , 58, 1855-1862	5	9
60	SLC10A1 S267F variant influences susceptibility to HBV infection and reduces cholesterol level by impairing bile acid uptake. <i>Journal of Viral Hepatitis</i> , 2019 , 26, 1178-1185	3.4	4
59	Systematic Functional Interrogation of Genes in GWAS Loci Identified ATF1 as a Key Driver in Colorectal Cancer Modulated by a Promoter-Enhancer Interaction. <i>American Journal of Human Genetics</i> , 2019 , 105, 29-47	11	26
58	A genetic variant in PIK3R1 is associated with pancreatic cancer survival in the Chinese population. <i>Cancer Medicine</i> , 2019 , 8, 3575-3582	4.8	7
57	Integrative analysis identifies genetic variant modulating MICA expression and altering susceptibility to persistent HBV infection. <i>Liver International</i> , 2019 , 39, 1927-1936	7.9	4
56	A missense variant in PTPN12 associated with the risk of colorectal cancer by modifying Ras/MEK/ERK signaling. <i>Cancer Epidemiology</i> , 2019 , 59, 109-114	2.8	7

55	Response to the comments on Status Matters: Relation between MHC levels and Hepatitis B persistence <i>Liver International</i> , 2019 , 39, 2208	7.9	
54	AWESOME: a database of SNPs that affect protein post-translational modifications. <i>Nucleic Acids Research</i> , 2019 , 47, D874-D880	20.1	32
53	A functional variant rs1537373 in 9p21.3 region is associated with pancreatic cancer risk. <i>Molecular Carcinogenesis</i> , 2019 , 58, 760-766	5	4
52	CancerSplicingQTL: a database for genome-wide identification of splicing QTLs in human cancer. <i>Nucleic Acids Research</i> , 2019 , 47, D909-D916	20.1	25
51	Exome-wide analyses identify low-frequency variant in CYP26B1 and additional coding variants associated with esophageal squamous cell carcinoma. <i>Nature Genetics</i> , 2018 , 50, 338-343	36.3	62
50	Integrative expression quantitative trait locus-based analysis of colorectal cancer identified a functional polymorphism regulating SLC22A5 expression. <i>European Journal of Cancer</i> , 2018 , 93, 1-9	7.5	40
49	Association of co-exposure to heavy metals with renal function in a hypertensive population. <i>Environment International</i> , 2018 , 112, 198-206	12.9	17
48	Educational and Behavioral Counseling in a Methadone Maintenance Treatment Program in China: A Randomized Controlled Trial. <i>Frontiers in Psychiatry</i> , 2018 , 9, 113	5	6
47	A Rare Missense Variant in TCF7L2 Associates with Colorectal Cancer Risk by Interacting with a GWAS-Identified Regulatory Variant in the MYC Enhancer. <i>Cancer Research</i> , 2018 , 78, 5164-5172	10.1	46
46	A Rare Variant P507L in TPP1 Interrupts TPP1-TIN2 Interaction, Influences Telomere Length, and Confers Colorectal Cancer Risk in Chinese Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 1029-1035	4	34
45	Associations of environmental exposure to metals with the risk of hypertension in China. <i>Science of the Total Environment</i> , 2018 , 622-623, 184-191	10.2	23
44	Integrative functional genomics identifies regulatory genetic variant modulating RAB31 expression and altering susceptibility to breast cancer. <i>Molecular Carcinogenesis</i> , 2018 , 57, 1845-1854	5	2
43	Exome-wide analysis identifies three low-frequency missense variants associated with pancreatic cancer risk in Chinese populations. <i>Nature Communications</i> , 2018 , 9, 3688	17.4	25
42	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. <i>Gut</i> , 2017 , 66, 581-587	19.2	51
41	A low-frequency variant in SMAD7 modulates TGF- β signaling and confers risk for colorectal cancer in Chinese population. <i>Molecular Carcinogenesis</i> , 2017 , 56, 1798-1807	5	43
40	Identification of a functional polymorphism affecting microRNA binding in the susceptibility locus 1q25.3 for colorectal cancer. <i>Molecular Carcinogenesis</i> , 2017 , 56, 2014-2021	5	7
39	Breast cancer risk-associated variants at 6q25.1 influence risk of hepatocellular carcinoma in a Chinese population. <i>Carcinogenesis</i> , 2017 , 38, 447-454	4.6	4
38	A functional variant in GREM1 confers risk for colorectal cancer by disrupting a hsa-miR-185-3p binding site. <i>Oncotarget</i> , 2017 , 8, 61318-61326	3.3	13

37	BRCA1 missense polymorphisms are associated with poor prognosis of pancreatic cancer patients in a Chinese population. <i>Oncotarget</i> , 2017 , 8, 36033-36039	3.3	19
36	A functional polymorphism located at transcription factor binding sites, rs6695837 near LAMC1 gene, confers risk of colorectal cancer in Chinese populations. <i>Carcinogenesis</i> , 2017 , 38, 177-183	4.6	54
35	Identification of a Potential Regulatory Variant for Colorectal Cancer Risk Mapping to 3p21.31 in Chinese Population. <i>Scientific Reports</i> , 2016 , 6, 25194	4.9	3
34	Association between bilirubin and risk of Non-Alcoholic Fatty Liver Disease based on a prospective cohort study. <i>Scientific Reports</i> , 2016 , 6, 31006	4.9	25
33	Genetic variant in DIP2A gene is associated with developmental dyslexia in Chinese population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016 , 171B, 203-8	3.5	23
32	Nighttime sleep duration and risk of nonalcoholic fatty liver disease: the Dongfeng-Tongji prospective study. <i>Annals of Medicine</i> , 2016 , 48, 468-476	1.5	11
31	A functional polymorphism in lnc-LAMC2-1:1 confers risk of colorectal cancer by affecting miRNA binding. <i>Carcinogenesis</i> , 2016 , 37, 443-51	4.6	60
30	The Roles of Genes in the Neuronal Migration and Neurite Outgrowth Network in Developmental Dyslexia: Single- and Multiple-Risk Genetic Variants. <i>Molecular Neurobiology</i> , 2016 , 53, 3967-3975	6.2	30
29	The contribution of serum hepatitis B virus load in the carcinogenesis and prognosis of hepatocellular carcinoma: evidence from two meta-analyses. <i>Oncotarget</i> , 2016 , 7, 49299-49309	3.3	9
28	A single nucleotide polymorphism in the 3'SUTR of STAT3 regulates its expression and reduces risk of pancreatic cancer in a Chinese population. <i>Oncotarget</i> , 2016 , 7, 62305-62311	3.3	8
27	A functional variant rs4442975 modulating FOXA1-binding affinity does not influence the risk or progression of breast cancer in Chinese Han population. <i>Oncotarget</i> , 2016 , 7, 81691-81697	3.3	3
26	Identification of a six microRNA signature as a novel potential prognostic biomarker in patients with head and neck squamous cell carcinoma. <i>Oncotarget</i> , 2016 , 7, 21579-90	3.3	25
25	Identification of a functional variant for colorectal cancer risk mapping to chromosome 5q31.1. <i>Oncotarget</i> , 2016 , 7, 35199-207	3.3	11
24	A novel subtype classification and risk of breast cancer by histone modification profiling. <i>Breast Cancer Research and Treatment</i> , 2016 , 157, 267-279	4.4	17
23	Genetic variants in the regulatory region of SLC10A1 are not associated with the risk of hepatitis B virus infection and clearance. <i>Infection, Genetics and Evolution</i> , 2016 , 44, 495-500	4.5	22
22	Genetic variants in the SWI/SNF complex and smoking collaborate to modify the risk of pancreatic cancer in a Chinese population. <i>Molecular Carcinogenesis</i> , 2015 , 54, 761-8	5	28
21	MAD1L1 Arg558His and MAD2L1 Leu84Met interaction with smoking increase the risk of colorectal cancer. <i>Scientific Reports</i> , 2015 , 5, 12202	4.9	14
20	Systematic confirmation study of GWAS-identified genetic variants for Kawasaki disease in a Chinese population. <i>Scientific Reports</i> , 2015 , 5, 8194	4.9	23

19	Dietary legume consumption reduces risk of colorectal cancer: evidence from a meta-analysis of cohort studies. <i>Scientific Reports</i> , 2015 , 5, 8797	4.9	58
18	A phosphorylation-related variant ADD1-rs4963 modifies the risk of colorectal cancer. <i>PLoS ONE</i> , 2015 , 10, e0121485	3.7	8
17	SF3A1 and pancreatic cancer: new evidence for the association of the spliceosome and cancer. <i>Oncotarget</i> , 2015 , 6, 37750-7	3.3	13
16	Genetic variant in SWI/SNF complexes influences hepatocellular carcinoma risk: a new clue for the contribution of chromatin remodeling in carcinogenesis. <i>Scientific Reports</i> , 2014 , 4, 4147	4.9	21
15	Variants in the 5Supstream region of GPC5 confer risk of lung cancer in never smokers. <i>Cancer Epidemiology</i> , 2014 , 38, 66-72	2.8	8
14	Allium vegetables and garlic supplements do not reduce risk of colorectal cancer, based on meta-analysis of prospective studies. <i>Clinical Gastroenterology and Hepatology</i> , 2014 , 12, 1991-2001.e1-4; quiz e121	6.9	41
13	Non-linear dose-response relationship between cigarette smoking and pancreatic cancer risk: evidence from a meta-analysis of 42 observational studies. <i>European Journal of Cancer</i> , 2014 , 50, 193-203	7.5	53
12	Parity and pancreatic cancer risk: evidence from a meta-analysis of twenty epidemiologic studies. <i>Scientific Reports</i> , 2014 , 4, 5313	4.9	18
11	8p22-23-rs2254546 as a susceptibility locus for Kawasaki disease: a case-control study and a meta-analysis. <i>Scientific Reports</i> , 2014 , 4, 4247	4.9	8
10	The roles of Ca ²⁺ /NFAT signaling genes in Kawasaki disease: single- and multiple-risk genetic variants. <i>Scientific Reports</i> , 2014 , 4, 5208	4.9	8
9	Genetic variant in MTRR, but not MTR, is associated with risk of congenital heart disease: an integrated meta-analysis. <i>PLoS ONE</i> , 2014 , 9, e89609	3.7	18
8	New integrated strategy emphasizing infection source control to curb Schistosomiasis japonica in a marshland area of Hubei Province, China: findings from an eight-year longitudinal survey. <i>PLoS ONE</i> , 2014 , 9, e89779	3.7	22
7	Dietary mushroom intake may reduce the risk of breast cancer: evidence from a meta-analysis of observational studies. <i>PLoS ONE</i> , 2014 , 9, e93437	3.7	31
6	A genetic variant rs1801274 in FCGR2A as a potential risk marker for Kawasaki disease: a case-control study and meta-analysis. <i>PLoS ONE</i> , 2014 , 9, e103329	3.7	26
5	Integrative genomic analysis identifies that SERPINA6-rs1998056 regulated by FOXA/ER α s associated with female hepatocellular carcinoma. <i>PLoS ONE</i> , 2014 , 9, e107246	3.7	8
4	Meta-analysis of the association between DCDC2 polymorphisms and risk of dyslexia. <i>Molecular Neurobiology</i> , 2013 , 47, 435-42	6.2	18
3	Genetic variations in the TGF β signaling pathway, smoking and risk of colorectal cancer in a Chinese population. <i>Carcinogenesis</i> , 2013 , 34, 936-42	4.6	67
2	Genetic variations in TERT-CLPTM1L locus are associated with risk of lung cancer in Chinese population. <i>Molecular Carcinogenesis</i> , 2013 , 52 Suppl 1, E118-26	5	42

- 1 HBV-related hepatocellular carcinoma susceptibility gene KIF1B is not associated with development of chronic hepatitis B. *PLoS ONE*, **2012**, 7, e28839 3·7 20