

# Xiaoping Miao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5659231/publications.pdf>

Version: 2024-02-01

286  
papers

20,413  
citations

16411

64  
h-index

13727

129  
g-index

293  
all docs

293  
docs citations

293  
times ranked

33156  
citing authors

#	ARTICLE	IF	CITATIONS
1	Neurologic Manifestations of Hospitalized Patients With Coronavirus Disease 2019 in Wuhan, China. <i>JAMA Neurology</i> , 2020, 77, 683.	4.5	5,308
2	Hirschsprung disease, associated syndromes and genetics: a review. <i>Journal of Medical Genetics</i> , 2007, 45, 1-14.	1.5	848
3	Clinical characteristics and risk factors associated with COVID-19 disease severity in patients with cancer in Wuhan, China: a multicentre, retrospective, cohort study. <i>Lancet Oncology</i> , The, 2020, 21, 893-903.	5.1	421
4	Skin damage among health care workers managing coronavirus disease-2019. <i>Journal of the American Academy of Dermatology</i> , 2020, 82, 1215-1216.	0.6	399
5	Genome-wide association study identifies three new susceptibility loci for esophageal squamous-cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2011, 43, 679-684.	9.4	260
6	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. <i>Nature Genetics</i> , 2011, 43, 1215-1218.	9.4	250
7	Genome-wide association analyses of esophageal squamous cell carcinoma in Chinese identify multiple susceptibility loci and gene-environment interactions. <i>Nature Genetics</i> , 2012, 44, 1090-1097.	9.4	238
8	Pancreatic cancer risk variant in LINC00673 creates a miR-1231 binding site and interferes with PTPN11 degradation. <i>Nature Genetics</i> , 2016, 48, 747-757.	9.4	237
9	Differential Contributions of Rare and Common, Coding and Noncoding Ret Mutations to Multifactorial Hirschsprung Disease Liability. <i>American Journal of Human Genetics</i> , 2010, 87, 60-74.	2.6	230
10	Cohort Profile: The Dongfengâ€Tongji cohort study of retired workers. <i>International Journal of Epidemiology</i> , 2013, 42, 731-740.	0.9	219
11	Metaâ€analysis on night shift work and risk of metabolic syndrome. <i>Obesity Reviews</i> , 2014, 15, 709-720.	3.1	218
12	Genetic polymorphisms of interleukin (IL)-1B, IL-1RN, IL-8, IL-10 and tumor necrosis factor Î± and risk of gastric cancer in a Chinese population. <i>Carcinogenesis</i> , 2005, 26, 631-636.	1.3	212
13	Identification of Genetic Variants in Base Excision Repair Pathway and Their Associations with Risk of Esophageal Squamous Cell Carcinoma. <i>Cancer Research</i> , 2004, 64, 4378-4384.	0.4	208
14	Clinical Characteristics of and Medical Interventions for COVID-19 in Hemodialysis Patients in Wuhan, China. <i>Journal of the American Society of Nephrology: JASN</i> , 2020, 31, 1387-1397.	3.0	206
15	lncRNASNP: a database of SNPs in lncRNAs and their potential functions in human and mouse. <i>Nucleic Acids Research</i> , 2015, 43, D181-D186.	6.5	204
16	Identification of Functional Genetic Variants in Cyclooxygenase-2 and Their Association With Risk of Esophageal Cancer. <i>Gastroenterology</i> , 2005, 129, 565-576.	0.6	200
17	PancanQTL: systematic identification of cis-eQTLs and trans-eQTLs in 33 cancer types. <i>Nucleic Acids Research</i> , 2018, 46, D971-D976.	6.5	191
18	Polymorphisms of Death Pathway Genes FAS and FASL in Esophageal Squamous-Cell Carcinoma. <i>Journal of the National Cancer Institute</i> , 2004, 96, 1030-1036.	3.0	185

#	ARTICLE	IF	CITATIONS
19	Functional Haplotypes in the Promoter of Matrix Metalloproteinase-2 Predict Risk of the Occurrence and Metastasis of Esophageal Cancer. <i>Cancer Research</i> , 2004, 64, 7622-7628.	0.4	182
20	Genome-wide association study identifies <i>NRG1</i> as a susceptibility locus for Hirschsprung's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 2694-2699.	3.3	171
21	The Role of P53 and MDM2 Polymorphisms in the Risk of Esophageal Squamous Cell Carcinoma. <i>Cancer Research</i> , 2005, 65, 9582-9587.	0.4	169
22	Genome-wide association study identifies five loci associated with susceptibility to pancreatic cancer in Chinese populations. <i>Nature Genetics</i> , 2012, 44, 62-66.	9.4	164
23	Clinical characteristics of 54 medical staff with COVID-19: A retrospective study in a single center in Wuhan, China. <i>Journal of Medical Virology</i> , 2020, 92, 807-813.	2.5	153
24	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2014, 46, 1001-1006.	9.4	148
25	Genetic polymorphisms in cell cycle regulatory genes MDM2 and TP53 are associated with susceptibility to lung cancer. <i>Human Mutation</i> , 2006, 27, 110-117.	1.1	142
26	Aurora-A/STK15 T + 91A is a general low penetrance cancer susceptibility gene: a meta-analysis of multiple cancer types. <i>Carcinogenesis</i> , 2005, 26, 1368-1373.	1.3	132
27	Polymorphisms in DNA base excision repair genes ADPR1 and XRCC1 and risk of lung cancer. <i>Cancer Research</i> , 2005, 65, 722-6.	0.4	127
28	Polymorphisms of DNA repair genes XRCC1 and XPD and their associations with risk of esophageal squamous cell carcinoma in a Chinese population. <i>International Journal of Cancer</i> , 2002, 100, 600-605.	2.3	120
29	Functional polymorphisms in cell death pathway genes FAS and FASL contribute to risk of lung cancer. <i>Journal of Medical Genetics</i> , 2005, 42, 479-484.	1.5	120
30	Genome-wide association study identifies a susceptibility locus for biliary atresia on 10q24.2. <i>Human Molecular Genetics</i> , 2010, 19, 2917-2925.	1.4	117
31	Urinary levels of bisphenol A, F and S and markers of oxidative stress among healthy adult men: Variability and association analysis. <i>Environment International</i> , 2019, 123, 301-309.	4.8	117
32	A novel T-77C polymorphism in DNA repair gene XRCC1 contributes to diminished promoter activity and increased risk of non-small cell lung cancer. <i>Oncogene</i> , 2006, 25, 3613-3620.	2.6	115
33	A genome-wide association study identifies two new cervical cancer susceptibility loci at 4q12 and 17q12. <i>Nature Genetics</i> , 2013, 45, 918-922.	9.4	108
34	Substantial reduction in risk of breast cancer associated with genetic polymorphisms in the promoters of the matrix metalloproteinase-2 and tissue inhibitor of metalloproteinase-2 genes. <i>Carcinogenesis</i> , 2003, 25, 399-404.	1.3	107
35	Meta-analysis of genome-wide association studies of adult height in East Asians identifies 17 novel loci. <i>Human Molecular Genetics</i> , 2015, 24, 1791-1800.	1.4	105
36	FASL -844C polymorphism is associated with increased activation-induced T cell death and risk of cervical cancer. <i>Journal of Experimental Medicine</i> , 2005, 202, 967-974.	4.2	104

#	ARTICLE	IF	CITATIONS
37	Sequence variations in the DNA repair gene XPD and risk of lung cancer in a Chinese population. <i>International Journal of Cancer</i> , 2003, 105, 669-673.	2.3	102
38	ELABELA and an ELABELA Fragment Protect against AKI. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 2694-2707.	3.0	101
39	Identification of Functional Genetic Variants in and Their Association With Risk of Esophageal Cancer. <i>Gastroenterology</i> , 2005, 129, 565-576.	0.6	100
40	The Expression Levels of Plasma microRNAs in Atrial Fibrillation Patients. <i>PLoS ONE</i> , 2012, 7, e44906.	1.1	97
41	Potential Influence of Menstrual Status and Sex Hormones on Female Severe Acute Respiratory Syndrome Coronavirus 2 Infection: A Cross-sectional Multicenter Study in Wuhan, China. <i>Clinical Infectious Diseases</i> , 2021, 72, e240-e248.	2.9	96
42	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	1.4	90
43	DNA repair gene XPC genotypes/haplotypes and risk of lung cancer in a Chinese population. <i>International Journal of Cancer</i> , 2005, 115, 478-483.	2.3	87
44	Analysis of Ovarian Injury Associated With COVID-19 Disease in Reproductive-Aged Women in Wuhan, China: An Observational Study. <i>Frontiers in Medicine</i> , 2021, 8, 635255.	1.2	86
45	Reelin gene variants and risk of autism spectrum disorders: An integrated meta-analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 192-200.	1.1	85
46	Common genetic variants regulating ADD3 gene expression alter biliary atresia risk. <i>Journal of Hepatology</i> , 2013, 59, 1285-1291.	1.8	84
47	Identification of a HOXD13 mutation in a VACTERL patient. <i>American Journal of Medical Genetics, Part A</i> , 2008, 146A, 3181-3185.	0.7	82
48	Functional Variants in Cell Death Pathway Genes and Risk of Pancreatic Cancer. <i>Clinical Cancer Research</i> , 2008, 14, 3230-3236.	3.2	82
49	Functional STK15 Phe31Ile Polymorphism Is Associated with the Occurrence and Advanced Disease Status of Esophageal Squamous Cell Carcinoma. <i>Cancer Research</i> , 2004, 64, 2680-2683.	0.4	80
50	Dietary legume consumption reduces risk of colorectal cancer: evidence from a meta-analysis of cohort studies. <i>Scientific Reports</i> , 2015, 5, 8797.	1.6	79
51	Functional haplotypes in the promoter of matrix metalloproteinase-2 and lung cancer susceptibility. <i>Carcinogenesis</i> , 2005, 26, 1117-1121.	1.3	78
52	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
53	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.	9.4	75
54	Genetic Variants in the Folate Pathway and the Risk of Neural Tube Defects: A Meta-Analysis of the Published Literature. <i>PLoS ONE</i> , 2013, 8, e59570.	1.1	74

#	ARTICLE	IF	CITATIONS
55	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016, 7, 11478.	5.8	73
56	A functional polymorphism in the matrix metalloproteinase-2 gene promoter (-1306C/T) is associated with risk of development but not metastasis of gastric cardia adenocarcinoma. <i>Cancer Research</i> , 2003, 63, 3987-90.	0.4	73
57	Association between serum uric acid and the metabolic syndrome among a middle- and old-age Chinese population. <i>European Journal of Epidemiology</i> , 2013, 28, 669-676.	2.5	72
58	Exposure to the Chinese Famine in Childhood Increases Type 2 Diabetes Risk in Adults. <i>Journal of Nutrition</i> , 2016, 146, 2289-2295.	1.3	70
59	Glyceraldehydeâ€³â€¢phosphate dehydrogenase promotes liver tumorigenesis by modulating phosphoglycerate dehydrogenase. <i>Hepatology</i> , 2017, 66, 631-645.	3.6	70
60	Environmental exposure to metals and the risk of hypertension: A cross-sectional study in China. <i>Environmental Pollution</i> , 2018, 233, 670-678.	3.7	70
61	Genetic variations in the TGFâ€¢ signaling pathway, smoking and risk of colorectal cancer in a Chinese population. <i>Carcinogenesis</i> , 2013, 34, 936-942.	1.3	69
62	Evaluation of Genetic Susceptibility Loci for Chronic Hepatitis B in Chinese: Two Independent Case-Control Studies. <i>PLoS ONE</i> , 2011, 6, e17608.	1.1	69
63	A common polymorphism in the 3â€¢UTR of cyclooxygenase 2/prostaglandin synthase 2 gene and risk of lung cancer in a Chinese population. <i>Lung Cancer</i> , 2005, 48, 11-17.	0.9	68
64	Interaction of P53 Arg72Pro and MDM2 T309G polymorphisms and their associations with risk of gastric cardia cancer. <i>Carcinogenesis</i> , 2007, 28, 1996-2001.	1.3	68
65	A functional polymorphism in <i>lnc-LAMC2-1:1</i> confers risk of colorectal cancer by affecting miRNA binding. <i>Carcinogenesis</i> , 2016, 37, 443-451.	1.3	68
66	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.	6.1	68
67	A genome wide association study of genetic loci that influence tumour biomarkers cancer antigen 19-9, carcinoembryonic antigen and Î± fetoprotein and their associations with cancer risk. <i>Gut</i> , 2014, 63, 143-151.	6.1	67
68	Longer Sleep Duration and Midday Napping Are Associated with a Higher Risk of CHD Incidence in Middle-Aged and Older Chinese: the Dongfeng-Tongji Cohort Study. <i>Sleep</i> , 2016, 39, 645-652.	0.6	64
69	Non-linear doseâ€¢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.	1.3	63
70	The effect of laparoscopic excision vs open excision in children with choledochal cyst: a midterm follow-up study. <i>Journal of Pediatric Surgery</i> , 2011, 46, 662-665.	0.8	61
71	Identification of Genes Associated With Hirschsprung Disease, Based on Whole-Genome Sequence Analysis, and Potential Effects on Enteric Nervous System Development. <i>Gastroenterology</i> , 2018, 155, 1908-1922.e5.	0.6	61
72	CancerSplicingQTL: a database for genome-wide identification of splicing QTLs in human cancer. <i>Nucleic Acids Research</i> , 2019, 47, D909-D916.	6.5	61

#	ARTICLE	IF	CITATIONS
73	A functional polymorphism located at transcription factor binding sites, rs6695837 near LAMC1 gene, confers risk of colorectal cancer in Chinese populations. <i>Carcinogenesis</i> , 2016, 38, bgw204.	1.3	59
74	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017, 152, 2011-2021.	0.6	58
75	Severe acute respiratory syndrome coronavirus 2 detection in the female lower genital tract. <i>American Journal of Obstetrics and Gynecology</i> , 2020, 223, 131-134.	0.7	58
76	Functional Phe31Ile polymorphism in Aurora A and risk of breast carcinoma. <i>Carcinogenesis</i> , 2004, 25, 2225-2230.	1.3	57
77	A Genome Wide Association Study Identifies Common Variants Associated with Lipid Levels in the Chinese Population. <i>PLoS ONE</i> , 2013, 8, e82420.	1.1	57
78	Genetic Polymorphisms in Methylenetetrahydrofolate Reductase and Thymidylate Synthase and Risk of Pancreatic Cancer. <i>Clinical Gastroenterology and Hepatology</i> , 2005, 3, 743-751.	2.4	56
79	Association between polymorphisms of <i>DRD2</i> and <i>DRD4</i> and opioid dependence: Evidence from the current studies. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011, 156, 661-670.	1.1	55
80	Prevalence and Associated Risk Factors of Dyslexic Children in a Middle-Sized City of China: A Cross-Sectional Study. <i>PLoS ONE</i> , 2013, 8, e56688.	1.1	55
81	Polymorphisms of <i>XRCC1</i> genes and risk of nasopharyngeal carcinoma in the Cantonese population. <i>BMC Cancer</i> , 2006, 6, 167.	1.1	54
82	A Rare Missense Variant in <i>TCF7L2</i> Associates with Colorectal Cancer Risk by Interacting with a GWAS-Identified Regulatory Variant in the <i>MYC</i> Enhancer. <i>Cancer Research</i> , 2018, 78, 5164-5172.	0.4	54
83	Inactivation of DNA repair gene O6-methylguanine-DNA methyltransferase by promoter hypermethylation and its relation to p53 mutations in esophageal squamous cell carcinoma. <i>Carcinogenesis</i> , 2003, 24, 1039-1044.	1.3	53
84	Adenosine Diphosphate Ribosyl Transferase and X-Ray Repair Cross-Complementing 1 Polymorphisms in Gastric Cardia Cancer. <i>Gastroenterology</i> , 2006, 131, 420-427.	0.6	53
85	Genome-Wide Interrogation Identifies <i>YAP1</i> Variants Associated with Survival of Small-Cell Lung Cancer Patients. <i>Cancer Research</i> , 2010, 70, 9721-9729.	0.4	53
86	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.e4.	2.4	53
87	AWESOME: a database of SNPs that affect protein post-translational modifications. <i>Nucleic Acids Research</i> , 2019, 47, D874-D880.	6.5	53
88	N <sup>6</sup> -methyladenosine mRNA methylation of <i>PIK3CB</i> regulates AKT signalling to promote PTEN-deficient pancreatic cancer progression. <i>Gut</i> , 2020, 69, 2180-2192.	6.1	52
89	A functional polymorphism in the <i>SULT1A1</i> gene (G638A) is associated with risk of lung cancer in relation to tobacco smoking. <i>Carcinogenesis</i> , 2003, 25, 773-778.	1.3	51
90	Genome-Wide Copy Number Analysis Uncovers a New HSCR Gene: <i>NRG3</i> . <i>PLoS Genetics</i> , 2012, 8, e1002687.	1.5	51

#	ARTICLE	IF	CITATIONS
91	Mutations in the NRG1 gene are associated with Hirschsprung disease. <i>Human Genetics</i> , 2012, 131, 67-76.	1.8	51
92	Risk SNP-Mediated Enhancer-Promoter Interaction Drives Colorectal Cancer through Both <i>FADS2</i> and <i>AP002754.2</i> . <i>Cancer Research</i> , 2020, 80, 1804-1818.	0.4	50
93	Perturbation of Hoxb5 Signaling in Vagal Neural Crests Down-Regulates Ret Leading to Intestinal Hypoganglionosis in Mice. <i>Gastroenterology</i> , 2008, 134, 1104-1115.	0.6	49
94	A polymorphic MYC response element in KBTBD11 influences colorectal cancer risk, especially in interaction with an MYC-regulated SNP rs6983267. <i>Annals of Oncology</i> , 2018, 29, 632-639.	0.6	49
95	A low-frequency variant in SMAD7 modulates TGF $\beta$ <sup>2</sup> signaling and confers risk for colorectal cancer in Chinese population. <i>Molecular Carcinogenesis</i> , 2017, 56, 1798-1807.	1.3	48
96	Reducing protein regulator of cytokinesis 1 as a prospective therapy for hepatocellular carcinoma. <i>Cell Death and Disease</i> , 2018, 9, 534.	2.7	48
97	Bidirectional association between nonalcoholic fatty liver disease and type 2 diabetes in Chinese population: Evidence from the Dongfeng-Tongji cohort study. <i>PLoS ONE</i> , 2017, 12, e0174291.	1.1	48
98	Genetic polymorphism in myeloperoxidase but not GSTM1 is associated with risk of lung squamous cell carcinoma in a Chinese population. <i>International Journal of Cancer</i> , 2002, 102, 275-279.	2.3	47
99	Dinucleotide polymorphism of p73 gene is associated with a reduced risk of lung cancer in a Chinese population. <i>International Journal of Cancer</i> , 2005, 114, 455-460.	2.3	47
100	Implications of Endocrine Gland-Derived Vascular Endothelial Growth Factor/Prokineticin-1 Signaling in Human Neuroblastoma Progression. <i>Clinical Cancer Research</i> , 2007, 13, 868-875.	3.2	47
101	Reduced RET expression in gut tissue of individuals carrying risk alleles of Hirschsprung's disease. <i>Human Molecular Genetics</i> , 2010, 19, 1461-1467.	1.4	47
102	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.	1.3	47
103	Urinary biomarkers of phthalates exposure and risks of thyroid cancer and benign nodule. <i>Journal of Hazardous Materials</i> , 2020, 383, 121189.	6.5	46
104	Association of candidate genetic variations with gastric cardia adenocarcinoma in Chinese population: a multiple interaction analysis. <i>Carcinogenesis</i> , 2011, 32, 336-342.	1.3	45
105	Pancan-meQTL: a database to systematically evaluate the effects of genetic variants on methylation in human cancer. <i>Nucleic Acids Research</i> , 2019, 47, D1066-D1072.	6.5	45
106	Genetic variations in <i>TERT</i> , <i>CLPTM</i> , <i>L</i> locus are associated with risk of lung cancer in chinese population. <i>Molecular Carcinogenesis</i> , 2013, 52, 118-126.	1.3	44
107	COVID-19 in Peritoneal Dialysis Patients. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2021, 16, 121-123.	2.2	44
108	<i>Helicobacter pylori</i> infection is associated with type 2 diabetes among a middle-aged and old-age Chinese population. <i>Diabetes/Metabolism Research and Reviews</i> , 2016, 32, 95-101.	1.7	43

#	ARTICLE	IF	CITATIONS
109	Functional evaluation of missense variations in the human MAD1L1 and MAD2L1 genes and their impact on susceptibility to lung cancer. <i>Journal of Medical Genetics</i> , 2010, 47, 616-622.	1.5	42
110	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.	3.9	42
111	Interactions between Genetic Variants in the Adiponectin, Adiponectin Receptor 1 and Environmental Factors on the Risk of Colorectal Cancer. <i>PLoS ONE</i> , 2011, 6, e27301.	1.1	41
112	Exposure to the Chinese famine in early life and hypertension prevalence risk in adults. <i>Journal of Hypertension</i> , 2017, 35, 63-68.	0.3	41
113	Association of co-exposure to heavy metals with renal function in a hypertensive population. <i>Environment International</i> , 2018, 112, 198-206.	4.8	41
114	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.	1.1	41
115	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.	2.6	41
116	Circulating essential metals and lung cancer: Risk assessment and potential molecular effects. <i>Environment International</i> , 2019, 127, 685-693.	4.8	41
117	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.	1.1	40
118	Comparative Evaluation of Participation and Diagnostic Yield of Colonoscopy vs Fecal Immunochemical Test vs Risk-Adapted Screening in Colorectal Cancer Screening: Interim Analysis of a Multicenter Randomized Controlled Trial (TARGET-C). <i>American Journal of Gastroenterology</i> , 2020, 115, 1264-1274.	0.2	40
119	Loss of myeloid-related proteins 8 and myeloid-related proteins 14 expression in human esophageal squamous cell carcinoma correlates with poor differentiation. <i>World Journal of Gastroenterology</i> , 2004, 10, 1093.	1.4	40
120	The SLC4A7 variant rs4973768 is associated with breast cancer risk: evidence from a case-control study and a meta-analysis. <i>Breast Cancer Research and Treatment</i> , 2012, 136, 847-857.	1.1	39
121	Association between bilirubin and risk of Non-Alcoholic Fatty Liver Disease based on a prospective cohort study. <i>Scientific Reports</i> , 2016, 6, 31006.	1.6	39
122	Potential of Arbidol for Post-exposure Prophylaxis of COVID-19 Transmission: A Preliminary Report of a Retrospective Cohort Study. <i>Current Medical Science</i> , 2020, 40, 480-485.	0.7	39
123	Combined Effect of Genetic Polymorphisms in P53, P73, and MDM2 on Non-small Cell Lung Cancer Survival. <i>Journal of Thoracic Oncology</i> , 2011, 6, 1793-1800.	0.5	38
124	Genetic variant in <i>KIAA0319</i> , but not in <i>DYX1C1</i> , is associated with risk of dyslexia: An integrated meta-analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 970-976.	1.1	38
125	Epigenomic landscape of 5-hydroxymethylcytosine reveals its transcriptional regulation of lncRNAs in colorectal cancer. <i>British Journal of Cancer</i> , 2017, 116, 658-668.	2.9	38
126	Using different anthropometric indices to assess prediction ability of type 2 diabetes in elderly population: a 5-year prospective study. <i>BMC Geriatrics</i> , 2018, 18, 218.	1.1	38

#	ARTICLE	IF	CITATIONS
127	Common genetic polymorphisms in the 5' flanking Region of the SULT1A1 gene: haplotypes and their association with platelet enzymatic activity. <i>Pharmacogenetics and Genomics</i> , 2005, 15, 465-473.	0.7	37
128	Significant increase in risk of gastroesophageal cancer is associated with interaction between promoter polymorphisms in thymidylate synthase and serum folate status. <i>Carcinogenesis</i> , 2005, 26, 1430-1435.	1.3	37
129	Risk prediction of esophageal squamous-cell carcinoma with common genetic variants and lifestyle factors in Chinese population. <i>Carcinogenesis</i> , 2013, 34, 1782-1786.	1.3	37
130	CYP2C19 polymorphism and clinical outcomes among patients of different races treated with clopidogrel: A systematic review and meta-analysis. <i>Journal of Huazhong University of Science and Technology [Medical Sciences]</i> , 2015, 35, 147-156.	1.0	37
131	Fine Mapping of the NRG1 Hirschsprung's Disease Locus. <i>PLoS ONE</i> , 2011, 6, e16181.	1.1	37
132	Different Physical Activity Subtypes and Risk of Metabolic Syndrome in Middle-Aged and Older Chinese People. <i>PLoS ONE</i> , 2013, 8, e53258.	1.1	36
133	Platelet 12-lipoxygenase Arg261Gln polymorphism: functional characterization and association with risk of esophageal squamous cell carcinoma in combination with COX-2 polymorphisms. <i>Pharmacogenetics and Genomics</i> , 2007, 17, 197-205.	0.7	35
134	A functional polymorphism (rs1607167) in the matrix metalloproteinase-1 promoter is associated with development and progression of lung cancer. <i>Cancer</i> , 2011, 117, 5172-5181.	2.0	35
135	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, 761-768.	1.3	35
136	Sleep Duration and Midday Napping with 5-Year Incidence and Reversion of Metabolic Syndrome in Middle-Aged and Older Chinese. <i>Sleep</i> , 2016, 39, 1911-1918.	0.6	35
137	ANKLE1 N6A-related variant is associated with colorectal cancer risk by maintaining the genomic stability. <i>International Journal of Cancer</i> , 2020, 146, 3281-3293.	2.3	35
138	Genetic variants in m6A modification genes are associated with esophageal squamous-cell carcinoma in the Chinese population. <i>Carcinogenesis</i> , 2020, 41, 761-768.	1.3	35
139	Liver Intestine-Cadherin (CDH17) Haplotype Is Associated with Increased Risk of Hepatocellular Carcinoma. <i>Clinical Cancer Research</i> , 2006, 12, 5248-5252.	3.2	34
140	A Solute Carrier Family 22 Member 3 Variant rs3088442 G>A Associated with Coronary Heart Disease Inhibits Lipopolysaccharide-induced Inflammatory Response. <i>Journal of Biological Chemistry</i> , 2015, 290, 5328-5340.	1.6	34
141	Green tea consumption is associated with reduced incident CHD and improved CHD-related biomarkers in the Dongfeng-Tongji cohort. <i>Scientific Reports</i> , 2016, 6, 24353.	1.6	34
142	Long sleep duration and afternoon napping are associated with higher risk of incident diabetes in middle-aged and older Chinese: the Dongfeng-Tongji cohort study. <i>Annals of Medicine</i> , 2016, 48, 216-223.	1.5	34
143	Susceptibility to gastric cardia adenocarcinoma and genetic polymorphisms in methylenetetrahydrofolate reductase in an at-risk Chinese population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002, 11, 1454-8.	1.1	34
144	MNX1 (HLXB9) mutations in Currarino patients. <i>Journal of Pediatric Surgery</i> , 2009, 44, 1892-1898.	0.8	33

#	ARTICLE	IF	CITATIONS
145	Serologic Detection of SARS-CoV-2 Infections in Hemodialysis Centers: A Multicenter Retrospective Study in Wuhan, China. <i>American Journal of Kidney Diseases</i> , 2020, 76, 490-499.e1.	2.1	33
146	A functional $\gamma$ T<sup>T</sup>C polymorphism in XRCC1 is associated with risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 479-487.	1.1	32
147	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.	1.1	32
148	Exome-wide analysis identifies three low-frequency missense variants associated with pancreatic cancer risk in Chinese populations. <i>Nature Communications</i> , 2018, 9, 3688.	5.8	32
149	Assessment of XPD Lys751Gln and XRCC1 T<sup>T</sup>C polymorphisms in advanced non-small-cell lung cancer patients treated with platinum-based chemotherapy. <i>Lung Cancer</i> , 2011, 73, 110-115.	0.9	31
150	Systematic Confirmation Study of GWAS-Identified Genetic Variants for Kawasaki Disease in A Chinese Population. <i>Scientific Reports</i> , 2015, 5, 8194.	1.6	31
151	Variations in <i>HSPA1B</i> at 6p21.3 Are Associated with Lung Cancer Risk and Prognosis in Chinese Populations. <i>Cancer Research</i> , 2011, 71, 7576-7586.	0.4	30
152	Exome Sequencing Identified NRG3 as a Novel Susceptible Gene of Hirschsprung's Disease in a Chinese Population. <i>Molecular Neurobiology</i> , 2013, 47, 957-966.	1.9	30
153	Functional Polymorphisms in FAS/FASL System Increase the Risk of Neuroblastoma in Chinese Population. <i>PLoS ONE</i> , 2013, 8, e71656.	1.1	30
154	Prenatal phthalate exposure, birth outcomes and DNA methylation of Alu and LINE-1 repetitive elements: A pilot study in China. <i>Chemosphere</i> , 2018, 206, 759-765.	4.2	30
155	GSK-3<sup>β</sup> Polymorphism Discriminates Bipolar Disorder and Schizophrenia: A Systematic Meta-Analysis. <i>Molecular Neurobiology</i> , 2013, 48, 404-411.	1.9	29
156	Independent and joint effects of moderate alcohol consumption and smoking on the risks of non-alcoholic fatty liver disease in elderly Chinese men. <i>PLoS ONE</i> , 2017, 12, e0181497.	1.1	28
157	Correlation Between Genetic Variations in Hox Clusters and Hirschsprung's Disease. <i>Annals of Human Genetics</i> , 2007, 71, 526-536.	0.3	27
158	Fine mapping of the 9q31 Hirschsprung's disease locus. <i>Human Genetics</i> , 2010, 127, 675-683.	1.8	27
159	Polymorphisms of death pathway genes FAS and FASL and risk of nasopharyngeal carcinoma. <i>Molecular Carcinogenesis</i> , 2010, 49, 944-950.	1.3	27
160	Application of a Hybrid Model for Predicting the Incidence of Tuberculosis in Hubei, China. <i>PLoS ONE</i> , 2013, 8, e80969.	1.1	27
161	RET Mutational Spectrum in Hirschsprung Disease: Evaluation of 601 Chinese Patients. <i>PLoS ONE</i> , 2011, 6, e28986.	1.1	26
162	The epidemiology and etiology of influenza-like illness in Chinese children from 2008 to 2010. <i>Journal of Medical Virology</i> , 2012, 84, 672-678.	2.5	26

#	ARTICLE	IF	CITATIONS
163	BMI, Waist Circumference and All-Cause Mortality in a Middle-Aged and Elderly Chinese Population. <i>Journal of Nutrition, Health and Aging</i> , 2018, 22, 975-981.	1.5	26
164	Multi-loci analysis reveals the importance of genetic variations in sensitivity of platinum-based chemotherapy in non-small-cell lung cancer. <i>Molecular Carcinogenesis</i> , 2013, 52, 923-931.	1.3	25
165	Identification of a novel polymorphism Arg290Gln of esophageal cancer related gene 1 ( ECRG1 ) and its related risk to esophageal squamous cell carcinoma. <i>Carcinogenesis</i> , 2006, 27, 798-802.	1.3	24
166	Meta-analysis of the Association Between DCDC2 Polymorphisms and Risk of Dyslexia. <i>Molecular Neurobiology</i> , 2013, 47, 435-442.	1.9	24
167	The Leptin Gene Family and Colorectal Cancer: Interaction with Smoking Behavior and Family History of Cancer. <i>PLoS ONE</i> , 2013, 8, e60777.	1.1	23
168	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.	1.1	23
169	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.	1.6	23
170	Role of RET and KIF1B gene polymorphisms in risk of Hirschsprung's disease in Chinese population. <i>Gut</i> , 2007, 56, 736-736.	6.1	22
171	Association of shift-work, daytime napping, and nighttime sleep with cancer incidence and cancer-caused mortality in Dongfeng-tongji cohort study. <i>Annals of Medicine</i> , 2016, 48, 641-651.	1.5	22
172	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.	6.5	22
173	Smoking and COX-2 Functional Polymorphisms Interact to Increase the Risk of Gastric Cardia Adenocarcinoma in Chinese Population. <i>PLoS ONE</i> , 2011, 6, e21894.	1.1	22
174	HBV-Related Hepatocellular Carcinoma Susceptibility Gene KIF1B Is Not Associated with Development of Chronic Hepatitis B. <i>PLoS ONE</i> , 2012, 7, e28839.	1.1	21
175	A Common SMAD7 Variant Is Associated with Risk of Colorectal Cancer: Evidence from a Case-Control Study and a Meta-Analysis. <i>PLoS ONE</i> , 2012, 7, e33318.	1.1	21
176	Genome-wide copy number variation study in anorectal malformations. <i>Human Molecular Genetics</i> , 2013, 22, 621-631.	1.4	21
177	Inhibition of kinesin family member 20B sensitizes hepatocellular carcinoma cell to microtubule-targeting agents by blocking cytokinesis. <i>Cancer Science</i> , 2018, 109, 3450-3460.	1.7	21
178	Metabolically healthy obesity increased diabetes incidence in a middle-aged and elderly Chinese population. <i>Diabetes/Metabolism Research and Reviews</i> , 2020, 36, e3202.	1.7	21
179	BRCA1 missense polymorphisms are associated with poor prognosis of pancreatic cancer patients in a Chinese population. <i>Oncotarget</i> , 2017, 8, 36033-36039.	0.8	21
180	MAD1L1 Arg558His and MAD2L1 Leu84Met interaction with smoking increase the risk of colorectal cancer. <i>Scientific Reports</i> , 2015, 5, 12202.	1.6	20

#	ARTICLE	IF	CITATIONS
181	Parity and pancreatic cancer risk: evidence from a meta-analysis of twenty epidemiologic studies. <i>Scientific Reports</i> , 2014, 4, 5313.	1.6	20
182	Serum bilirubin levels and risk of type 2 diabetes: results from two independent cohorts in middle-aged and elderly Chinese. <i>Scientific Reports</i> , 2017, 7, 41338.	1.6	20
183	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.	0.8	20
184	SF3A1 and pancreatic cancer: new evidence for the association of the spliceosome and cancer. <i>Oncotarget</i> , 2015, 6, 37750-37757.	0.8	20
185	Haplotype Analysis Reveals a Possible Founder Effect of RET Mutation R114H for Hirschsprung's Disease in the Chinese Population. <i>PLoS ONE</i> , 2010, 5, e10918.	1.1	19
186	Identification of common variants in BRCA2 and MAP2K4 for susceptibility to sporadic pancreatic cancer. <i>Carcinogenesis</i> , 2013, 34, 1001-1005.	1.3	19
187	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	19
188	Association of regular physical activity with total and cause-specific mortality among middle-aged and older Chinese: a prospective cohort study. <i>Scientific Reports</i> , 2017, 7, 39939.	1.6	19
189	Interaction of <i>Cyclooxygenase-2</i> promoter polymorphisms with <i>Helicobacter pylori</i> infection and risk of gastric cancer. <i>Molecular Carcinogenesis</i> , 2011, 50, 876-883.	1.3	18
190	Risk of GWAS-identified genetic variants for breast cancer in a Chinese population: a multiple interaction analysis. <i>Breast Cancer Research and Treatment</i> , 2013, 142, 637-644.	1.1	18
191	Victims of Chinese famine in early life have increased risk of metabolic syndrome in adulthood. <i>Nutrition</i> , 2018, 53, 20-25.	1.1	18
192	Involvement of visinin-like protein-1 (VSNL-1) in regulating proliferative and invasive properties of neuroblastoma. <i>Carcinogenesis</i> , 2007, 28, 2122-2130.	1.3	17
193	Mutational analysis of <i>SHH</i> and <i>GLI3</i> in anorectal malformations. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2008, 82, 644-648.	1.6	17
194	Replication Study in Chinese Population and Meta-Analysis Supports Association of the 5p15.33 Locus with Lung Cancer. <i>PLoS ONE</i> , 2013, 8, e62485.	1.1	17
195	Repurposed Tocilizumab in Patients with Severe COVID-19. <i>Journal of Immunology</i> , 2021, 206, 599-606.	0.4	17
196	Identification of genetic variants in m6A modification genes associated with pancreatic cancer risk in the Chinese population. <i>Archives of Toxicology</i> , 2021, 95, 1117-1128.	1.9	17
197	Genetic Polymorphisms in Folate- Metabolizing Enzymes and Risk of Gastroesophageal Cancers: A Potential Nutrient-Gene Interaction in Cancer Development. <i>Forum of Nutrition</i> , 2007, 60, 140-145.	3.7	16
198	Mapping of a Hirschsprung's disease locus in 3p21. <i>European Journal of Human Genetics</i> , 2008, 16, 833-840.	1.4	16

#	ARTICLE	IF	CITATIONS
199	Polymorphisms of methylenetetrahydrofolate reductase are associated with a high risk of nasopharyngeal carcinoma in a smoking population from southern China. <i>Molecular Carcinogenesis</i> , 2010, 49, 928-934.	1.3	16
200	Association between serum bilirubin levels and decline in estimated glomerular filtration rate among patients with type 2 diabetes. <i>Journal of Diabetes and Its Complications</i> , 2016, 30, 1255-1260.	1.2	16
201	Polymorphisms in TP53 and MDM2 contribute to higher risk of colorectal cancer in Chinese population: a hospital-based, case-control study. <i>Molecular Biology Reports</i> , 2012, 39, 9661-9668.	1.0	15
202	Oncogenic B-RafV600E abrogates the AKT/B-Raf/Mps1 interaction in melanoma cells. <i>Cancer Letters</i> , 2013, 337, 125-132.	3.2	15
203	The Roles of Ca <sup>2+</sup> /NFAT Signaling Genes in Kawasaki Disease: Single- and Multiple-Risk Genetic Variants. <i>Scientific Reports</i> , 2014, 4, 5208.	1.6	15
204	Bidirectional association between nonalcoholic fatty liver disease and hypertension from the Dongfeng-Tongji cohort study. <i>Journal of the American Society of Hypertension</i> , 2018, 12, 660-670.	2.3	15
205	The SNP rs961253 in 20p12.3 Is Associated with Colorectal Cancer Risk: A Case-Control Study and a Meta-Analysis of the Published Literature. <i>PLoS ONE</i> , 2012, 7, e34625.	1.1	14
206	Associations between Two Genetic Variants in NKX2-5 and Risk of Congenital Heart Disease in Chinese Population: A Meta-Analysis. <i>PLoS ONE</i> , 2013, 8, e70979.	1.1	14
207	A genetic variant in microRNA target site of TGF- $\beta$ <sup>2</sup> signaling pathway increases the risk of colorectal cancer in a Chinese population. <i>Tumor Biology</i> , 2014, 35, 4301-4306.	0.8	14
208	Serum bilirubin concentrations and incident coronary heart disease risk among patients with type 2 diabetes: the Dongfeng-Tongji cohort. <i>Acta Diabetologica</i> , 2017, 54, 257-264.	1.2	14
209	Aberrant RNA Splicing Is a Primary Link between Genetic Variation and Pancreatic Cancer Risk. <i>Cancer Research</i> , 2022, 82, 2084-2096.	0.4	14
210	Evaluation of the Thyroid Transcription Factor-1 Gene (TITF1) as a Hirschsprung's Disease Locus. <i>Annals of Human Genetics</i> , 2007, 71, 746-754.	0.3	13
211	Serum creatinine levels and risk of metabolic syndrome in a middle-aged and older Chinese population. <i>Clinica Chimica Acta</i> , 2015, 440, 177-182.	0.5	13
212	Effect modification of CPY2E1 and GSTZ1 genetic polymorphisms on associations between prenatal disinfection by-products exposure and birth outcomes. <i>Environmental Pollution</i> , 2018, 243, 1126-1133.	3.7	13
213	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.	1.3	13
214	LINC01149 variant modulates MICA expression that facilitates hepatitis B virus spontaneous recovery but increases hepatocellular carcinoma risk. <i>Oncogene</i> , 2020, 39, 1944-1956.	2.6	13
215	The SNP rs402710 in 5p15.33 Is Associated with Lung Cancer Risk: A Replication Study in Chinese Population and a Meta-Analysis. <i>PLoS ONE</i> , 2013, 8, e76252.	1.1	13
216	No association betweenXRCC1polymorphisms and survival in non-small-cell lung cancer patients treated with platinum-based chemotherapy. <i>Cancer Biology and Therapy</i> , 2010, 10, 854-859.	1.5	12

#	ARTICLE	IF	CITATIONS
217	Two Novel Variants on 13q22.1 Are Associated with Risk of Esophageal Squamous Cell Carcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1774-1780.	1.1	12
218	The Relationship between Serum Bilirubin and Elevated Fibrotic Indices among HBV Carriers: A Cross-Sectional Study of a Chinese Population. <i>International Journal of Molecular Sciences</i> , 2016, 17, 2057.	1.8	12
219	Association between resting heart rate and incident diabetes risk: a Mendelian randomization study. <i>Acta Diabetologica</i> , 2019, 56, 1037-1044.	1.2	12
220	Identification of a functional variant for colorectal cancer risk mapping to chromosome 5q31.1. <i>Oncotarget</i> , 2016, 7, 35199-35207.	0.8	12
221	Variants in the 5' upstream region of GPC5 confer risk of lung cancer in never smokers. <i>Cancer Epidemiology</i> , 2014, 38, 66-72.	0.8	11
222	8p22 rs2254546 as a Susceptibility Locus for Kawasaki Disease: a Case-control Study and a Meta-analysis. <i>Scientific Reports</i> , 2014, 4, 4247.	1.6	11
223	Bisphenol A exposure, interaction with genetic variants and colorectal cancer via mediating oxidative stress biomarkers. <i>Environmental Pollution</i> , 2021, 287, 117630.	3.7	11
224	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.	3.9	11
225	Lack of association between nNOS rs84G>A polymorphism and risk of infantile hypertrophic pyloric stenosis in a Chinese population. <i>Journal of Pediatric Surgery</i> , 2010, 45, 709-713.	0.8	10
226	Associations between daily cooking duration and the prevalence of diabetes and prediabetes in a middle-aged and elderly Chinese population: A cross-sectional study. <i>Indoor Air</i> , 2018, 28, 238-246.	2.0	10
227	Serum alanine transaminase levels predict type 2 diabetes risk among a middle-aged and elderly Chinese population. <i>Annals of Hepatology</i> , 2019, 18, 298-303.	0.6	10
228	Prediction of 5-year risk of diabetes mellitus in relatively low risk middle-aged and elderly adults. <i>Acta Diabetologica</i> , 2020, 57, 63-70.	1.2	10
229	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.	1.9	10
230	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.	1.3	10
231	A single nucleotide polymorphism in the 3' UTR of STAT3 regulates its expression and reduces risk of pancreatic cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 62305-62311.	0.8	10
232	Hepatocellular carcinoma risk variant modulates lncRNA HLA-DQB1-AS1 expression via a long-range enhancer-promoter interaction. <i>Carcinogenesis</i> , 2021, 42, 1347-1356.	1.3	10
233	Variants of HSPA1A in Combination with Plasma Hsp70 and Anti-Hsp70 Antibody Levels Associated with Higher Risk of Acute Coronary Syndrome. <i>Cardiology</i> , 2011, 119, 57-64.	0.6	9
234	Integrative Genomic Analysis Identifies That SERPINA6-rs1998056 Regulated by FOXA/ER Is Associated with Female Hepatocellular Carcinoma. <i>PLoS ONE</i> , 2014, 9, e107246.	1.1	9

#	ARTICLE	IF	CITATIONS
235	Association between polymorphisms within the susceptibility region 8q24 and breast cancer in a Chinese population. <i>Tumor Biology</i> , 2014, 35, 2649-2654.	0.8	9
236	Association of Adiposity Indices with Platelet Distribution Width and Mean Platelet Volume in Chinese Adults. <i>PLoS ONE</i> , 2015, 10, e0129677.	1.1	9
237	Development of a new scoring system to predict 5-year incident diabetes risk in middle-aged and older Chinese. <i>Acta Diabetologica</i> , 2018, 55, 13-19.	1.2	9
238	Educational and Behavioral Counseling in a Methadone Maintenance Treatment Program in China: A Randomized Controlled Trial. <i>Frontiers in Psychiatry</i> , 2018, 9, 113.	1.3	9
239	A genetic variant in <i>PIK3R1</i> is associated with pancreatic cancer survival in the Chinese population. <i>Cancer Medicine</i> , 2019, 8, 3575-3582.	1.3	9
240	A missense variant in <i>PTPN12</i> associated with the risk of colorectal cancer by modifying Ras/MEK/ERK signaling. <i>Cancer Epidemiology</i> , 2019, 59, 109-114.	0.8	9
241	A functional variant rs1537373 in 9p21.3 region is associated with pancreatic cancer risk. <i>Molecular Carcinogenesis</i> , 2019, 58, 760-766.	1.3	9
242	Efficacy of hydrogel patches in preventing facial skin damage caused by mask compression in fighting against coronavirus disease 2019: a short-term, self-controlled study. <i>Journal of the European Academy of Dermatology and Venereology</i> , 2020, 34, e441-e443.	1.3	9
243	Functional characterization of a low-frequency V1937I variant in <i>FASN</i> associated with susceptibility to esophageal squamous cell carcinoma. <i>Archives of Toxicology</i> , 2020, 94, 2039-2046.	1.9	9
244	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.	3.2	9
245	An Integrated Meta-Analysis of Two Variants in <i>HOXA1/HOXB1</i> and Their Effect on the Risk of Autism Spectrum Disorders. <i>PLoS ONE</i> , 2011, 6, e25603.	1.1	9
246	The Genetic Variant on Chromosome 10p14 Is Associated with Risk of Colorectal Cancer: Results from a Case-Control Study and a Meta-Analysis. <i>PLoS ONE</i> , 2013, 8, e64310.	1.1	9
247	A Functional Variant rs1820453 in <i>YAP1</i> and Breast Cancer Risk in Chinese Population. <i>PLoS ONE</i> , 2013, 8, e79056.	1.1	9
248	The Case-Only Test for Gene-Environment Interaction is Not Uniformly Powerful: An Empirical Example. <i>Genetic Epidemiology</i> , 2013, 37, 402-407.	0.6	8
249	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.	1.7	8
250	Urinary bisphenol A and its interaction with <i>CYP17A1</i> rs743572 are associated with breast cancer risk. <i>Chemosphere</i> , 2022, 286, 131880.	4.2	8
251	Genetic Predisposition to Colon and Rectal Adenocarcinoma Is Mediated by a Super-enhancer Polymorphism Coactivating <i>CD9</i> and <i>PLEKHG6</i> . <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 850-859.	1.1	8
252	MNS16A Tandem Repeats Minisatellite of Human Telomerase Gene and Cancer Risk: A Meta-Analysis. <i>PLoS ONE</i> , 2013, 8, e73367.	1.1	8

#	ARTICLE	IF	CITATIONS
253	A Phosphorylation-Related Variant ADD1-rs4963 Modifies the Risk of Colorectal Cancer. <i>PLoS ONE</i> , 2015, 10, e0121485.	1.1	8
254	Association of IL-1 Receptor Antagonist Gene VNTR Polymorphism with Ischemic Stroke in the Chinese Uyghur Population. <i>Biochemical Genetics</i> , 2013, 51, 698-706.	0.8	7
255	Healthy lifestyle and cancer risk among Chinese population in the Dongfeng-Tongji cohort. <i>Annals of Medicine</i> , 2020, 52, 393-402.	1.5	7
256	Challenges and recommendations for cancer care in the COVID-19 pandemic. <i>Cancer Biology and Medicine</i> , 2020, 17, 515-518.	1.4	7
257	A functional variant in TNXB promoter associates with the risk of esophageal squamous cell carcinoma. <i>Molecular Carcinogenesis</i> , 2020, 59, 439-446.	1.3	7
258	Genetic variants associated with expression of TCF19 contribute to the risk of head and neck cancer in Chinese population. <i>Journal of Medical Genetics</i> , 2021, , jmedgenet-2020-107410.	1.5	7
259	Leveraging Fecal Microbial Markers to Improve the Diagnostic Accuracy of the Fecal Immunochemical Test for Advanced Colorectal Adenoma. <i>Clinical and Translational Gastroenterology</i> , 2021, 12, e00389.	1.3	7
260	Genetic correction improves prediction efficiency of serum tumor biomarkers on digestive cancer risk in the elderly Chinese cohort study. <i>Oncotarget</i> , 2018, 9, 7389-7397.	0.8	7
261	Application of HapMap data to the evaluation of 8 candidate genes for pediatric slow transit constipation. <i>Journal of Pediatric Surgery</i> , 2007, 42, 666-671.	0.8	6
262	Housing Characteristics in Relation to Exhaled Nitric Oxide in China. <i>American Journal of Health Behavior</i> , 2015, 39, 88-98.	0.6	6
263	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.	1.6	6
264	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.	3.7	6
265	Association between polymorphisms in cdc27 and breast cancer in a Chinese population. <i>Tumor Biology</i> , 2015, 36, 5299-5304.	0.8	5
266	Disparities of time trends and birth cohort effects on invasive breast cancer incidence in Shanghai and Hong Kong pre- and post-menopausal women. <i>BMC Cancer</i> , 2017, 17, 362.	1.1	5
267	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, 1-10.	1.9	5
268	Integrated genome-wide gene expression map and high-resolution analysis of aberrant chromosomal regions in squamous cell lung cancer. <i>FEBS Letters</i> , 2006, 580, 2774-2778.	1.3	4
269	Allele-Specific Expression at the <i>RET</i> Locus in Blood and Gut Tissue of Individuals Carrying Risk Alleles for Hirschsprung Disease. <i>Human Mutation</i> , 2013, 34, 754-762.	1.1	4
270	A Functional p53 Responsive Polymorphism in KITLG, rs4590952, Does not Affect the Risk of Breast Cancer. <i>Scientific Reports</i> , 2014, 4, 6371.	1.6	4

#	ARTICLE	IF	CITATIONS
271	The advances of genetics research on Hirschsprung's disease. <i>Pediatric Investigation</i> , 2018, 2, 189-195.	0.6	4
272	Implications of Lifestyle Factors and Polygenic Risk Score for Absolute Risk Prediction of Colorectal Neoplasm and Risk-Adapted Screening. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 685410.	1.6	4
273	Inverse association between plasma homocysteine concentrations and type 2 diabetes mellitus among a middle-aged and elderly Chinese population. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2018, 28, 278-284.	1.1	3
274	Genetic correction of serum $\alpha$ -fetoprotein level improves risk prediction of primary hepatocellular carcinoma in the Dongfeng-Tongji cohort study. <i>Cancer Medicine</i> , 2018, 7, 2691-2698.	1.3	3
275	Trans-acting non-synonymous variant of FOXA1 predisposes to hepatocellular carcinoma through modulating FOXA1-ER $\alpha$ transcriptional program and may have undergone natural selection. <i>Carcinogenesis</i> , 2020, 41, 146-158.	1.3	3
276	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.	0.8	3
277	Safety of biologics for psoriasis patients during the COVID-19 pandemic: the experience from Wuhan, China. <i>European Journal of Dermatology</i> , 2020, 30, 738-740.	0.3	3
278	Systematic analysis on expression quantitative trait loci identifies a novel regulatory variant in ring finger and WD repeat domain 3 associated with prognosis of pancreatic cancer. <i>Chinese Medical Journal</i> , 2022, 135, 1348-1357.	0.9	3
279	ICPS: an integrative cancer profiler system. <i>Bioinformatics</i> , 2010, 26, 2649-2650.	1.8	2
280	No NRG1 V266L in Chinese patients with schizophrenia. <i>Psychiatric Genetics</i> , 2011, 21, 47-49.	0.6	2
281	Reply to comment: Serum bilirubin concentrations, type 2 diabetes, and incident coronary heart disease. <i>Acta Diabetologica</i> , 2019, 56, 383-384.	1.2	2
282	Association of blood pressure and long-term change with chronic kidney disease risk among Chinese adults with different glucose metabolism according to the 2017 ACC/AHA guidelines. <i>Journal of Clinical Hypertension</i> , 2021, , .	1.0	2
283	A novel variant on chromosome 6p21.1 is associated with the risk of developing colorectal cancer: a two-stage case-control study in Han Chinese. <i>BMC Cancer</i> , 2016, 16, 807.	1.1	1
284	Challenges and countermeasures in the prevention of nosocomial infections of SARS-CoV-2 before resumption of work: Implications for the dermatology department. <i>Journal of the American Academy of Dermatology</i> , 2020, 83, 961-963.	0.6	1
285	Polymorphisms and mutations of the folate receptor-alpha gene and risk of gastric cancer in a Chinese population. <i>International Journal of Molecular Medicine</i> , 2005, 15, 627.	1.8	0
286	FOXA1 of regulatory variant associated with risk of breast cancer through allele-specific enhancer in the Chinese population. <i>Breast Cancer</i> , 2021, , 1.	1.3	0