

Xiaoping Miao

List of Publications by Year in descending order

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Version: 2024-02-01

286
papers

20,413
citations

18887

64
h-index

15698

129
g-index

293
all docs

293
docs citations

293
times ranked

35393
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

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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

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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

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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

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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 ⁶ -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

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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 β ² 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> , and <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 Chinese population. <i>Diabetes/Metabolism Research and Reviews</i> , 2016, 32, 95-101.	1.7	43

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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 TPP1â€“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 caseâ€“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

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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 (rs16071G>2G) 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 N6⁶-Methyladenosine-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
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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