

# Meilin Wang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4787773/publications.pdf>

Version: 2024-02-01

179  
papers

5,721  
citations

81743

39  
h-index

110170

64  
g-index

182  
all docs

182  
docs citations

182  
times ranked

8395  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	LncRNA MT1JP functions as a ceRNA in regulating FBXW7 through competitively binding to miR-92a-3p in gastric cancer. <i>Molecular Cancer</i> , 2018, 17, 87.	7.9	218
3	Exosome-transmitted long non-coding RNA PTENP1 suppresses bladder cancer progression. <i>Molecular Cancer</i> , 2018, 17, 143.	7.9	217
4	Circular RNAs in body fluids as cancer biomarkers: the new frontier of liquid biopsies. <i>Molecular Cancer</i> , 2021, 20, 13.	7.9	176
5	Genome-wide association study identifies a new susceptibility locus for cleft lip with or without a cleft palate. <i>Nature Communications</i> , 2015, 6, 6414.	5.8	167
6	Genetic variants in lncRNA HOTAIR are associated with risk of colorectal cancer. <i>Mutagenesis</i> , 2015, 30, 303-310.	1.0	128
7	Multiomics Evaluation of Gastrointestinal and Other Clinical Characteristics of COVID-19. <i>Gastroenterology</i> , 2020, 158, 2298-2301.e7.	0.6	117
8	Identification of novel piRNAs in bladder cancer. <i>Cancer Letters</i> , 2015, 356, 561-567.	3.2	115
9	A functional polymorphism in <i>MSMB</i> gene promoter is associated with prostate cancer risk and serum MSMB expression. <i>Prostate</i> , 2010, 70, 1146-1152.	1.2	106
10	Circulating miR-497 and miR-663b in plasma are potential novel biomarkers for bladder cancer. <i>Scientific Reports</i> , 2015, 5, 10437.	1.6	105
11	The association analysis of lncRNA <i>HOTAIR</i> genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2015, 6, 31255-31262.	0.8	95
12	Polymorphism of the pre-miR-146a is associated with risk of cervical cancer in a Chinese population. <i>Gynecologic Oncology</i> , 2011, 122, 33-37.	0.6	92
13	Association of genetic variants in lncRNA <i>H19</i> with risk of colorectal cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 25470-25477.	0.8	90
14	Genome-wide association analysis of Vogt-Koyanagi-Harada syndrome identifies two new susceptibility loci at 1p31.2 and 10q21.3. <i>Nature Genetics</i> , 2014, 46, 1007-1011.	9.4	88
15	Genetic Variants in miRNAs Predict Bladder Cancer Risk and Recurrence. <i>Cancer Research</i> , 2012, 72, 6173-6182.	0.4	86
16	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016, 7, 11478.	5.8	73
17	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-elf3h interaction. <i>Molecular Cancer</i> , 2022, 21, 49.	7.9	72
18	Genetic variants in lncRNA <i>H19</i> are associated with the risk of bladder cancer in a Chinese population. <i>Mutagenesis</i> , 2016, 31, 531-538.	1.0	70

#	ARTICLE	IF	CITATIONS
19	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
20	Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. <i>Gene</i> , 2015, 556, 227-234.	1.0	66
21	A Functional Polymorphism in <i>miRNA-196a2</i> Is Associated with Colorectal Cancer Risk in a Chinese Population. <i>DNA and Cell Biology</i> , 2012, 31, 350-354.	0.9	63
22	Meta-analysis on the effectiveness of team-based learning on medical education in China. <i>BMC Medical Education</i> , 2018, 18, 77.	1.0	63
23	A Functional Polymorphism in <i>Pre-miR-146a</i> Is Associated with Susceptibility to Gastric Cancer in a Chinese Population. <i>DNA and Cell Biology</i> , 2012, 31, 1290-1295.	0.9	59
24	A genetic variant in <i>miR-146a</i> modifies colorectal cancer susceptibility in a Chinese population. <i>Archives of Toxicology</i> , 2013, 87, 825-833.	1.9	58
25	Global gene expression profiling of human bronchial epithelial cells exposed to airborne fine particulate matter collected from Wuhan, China. <i>Toxicology Letters</i> , 2014, 228, 25-33.	0.4	58
26	Exome Array Analysis Identifies Variants in <i>SPOCD1</i> and <i>BTN3A2</i> That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017, 152, 2011-2021.	0.6	58
27	Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. <i>Carcinogenesis</i> , 2014, 35, 2723-2730.	1.3	57
28	A novel antisense long noncoding RNA regulates the expression of <i>MDC1</i> in bladder cancer. <i>Oncotarget</i> , 2015, 6, 484-493.	0.8	56
29	Genetic variants in noncoding <i>PIWI</i> -interacting RNA and colorectal cancer risk. <i>Cancer</i> , 2015, 121, 2044-2052.	2.0	56
30	The classic EDCs, phthalate esters and organochlorines, in relation to abnormal sperm quality: a systematic review with meta-analysis. <i>Scientific Reports</i> , 2016, 6, 19982.	1.6	54
31	Genetic variant in <i>PSCA</i> predicts survival of diffuse-type gastric cancer in a Chinese population. <i>International Journal of Cancer</i> , 2011, 129, 1207-1213.	2.3	52
32	Folic acid supplements and colorectal cancer risk: meta-analysis of randomized controlled trials. <i>Scientific Reports</i> , 2015, 5, 12044.	1.6	51
33	Large-scale association analysis in Asians identifies new susceptibility loci for prostate cancer. <i>Nature Communications</i> , 2015, 6, 8469.	5.8	51
34	<i>miR-107</i> regulates tumor progression by targeting <i>NF1</i> in gastric cancer. <i>Scientific Reports</i> , 2016, 6, 36531.	1.6	51
35	Association of three polymorphisms in <i>ARID5B</i> , <i>IKZF1</i> and <i>CEBPE</i> with the risk of childhood acute lymphoblastic leukemia in a Chinese population. <i>Gene</i> , 2013, 524, 203-207.	1.0	47
36	Genetic variations in microRNAs and the risk and survival of renal cell cancer. <i>Carcinogenesis</i> , 2014, 35, 1629-1635.	1.3	47

#	ARTICLE	IF	CITATIONS
37	Genome-Wide Association Study of Bladder Cancer in a Chinese Cohort Reveals a New Susceptibility Locus at 5q12.3. <i>Cancer Research</i> , 2016, 76, 3277-3284.	0.4	46
38	Clinical significance of <i>SOD2</i> and <i>GSTP1</i> gene polymorphisms in Chinese patients with gastric cancer. <i>Cancer</i> , 2012, 118, 5489-5496.	2.0	43
39	A functional variant in miR-143 promoter contributes to prostate cancer risk. <i>Archives of Toxicology</i> , 2016, 90, 403-414.	1.9	43
40	An inverse association between tea consumption and colorectal cancer risk. <i>Oncotarget</i> , 2017, 8, 37367-37376.	0.8	42
41	KCNMA1 cooperating with PTK2 is a novel tumor suppressor in gastric cancer and is associated with disease outcome. <i>Molecular Cancer</i> , 2017, 16, 46.	7.9	41
42	Short-term effects of ambient air pollution and childhood lower respiratory diseases. <i>Scientific Reports</i> , 2017, 7, 4414.	1.6	41
43	Common genetic variants in pre-microRNAs are associated with risk of coal workers' pneumoconiosis. <i>Journal of Human Genetics</i> , 2010, 55, 13-17.	1.1	40
44	A Novel Functional Polymorphism C1797G in the MDM2 Promoter Is Associated with Risk of Bladder Cancer in a Chinese Population. <i>Clinical Cancer Research</i> , 2008, 14, 3633-3640.	3.2	39
45	Replication and cumulative effects of GWAS-identified genetic variations for prostate cancer in Asians: a case-control study in the ChinaPca consortium. <i>Carcinogenesis</i> , 2012, 33, 356-360.	1.3	38
46	Environmental factors, seven GWAS-identified susceptibility loci, and risk of gastric cancer and its precursors in a Chinese population. <i>Cancer Medicine</i> , 2017, 6, 708-720.	1.3	38
47	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020, 41, 8-17.	1.3	38
48	Environmental exposure to BDE47 is associated with increased diabetes prevalence: Evidence from community-based case-control studies and an animal experiment. <i>Scientific Reports</i> , 2016, 6, 27854.	1.6	37
49	Genome-wide long non-coding RNAs identified a panel of novel plasma biomarkers for gastric cancer diagnosis. <i>Gastric Cancer</i> , 2019, 22, 731-741.	2.7	37
50	The HOTAIR, PRNCR1 and POLR2E polymorphisms are associated with cancer risk: a meta-analysis. <i>Oncotarget</i> , 2017, 8, 43271-43283.	0.8	37
51	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. <i>Gut</i> , 2020, 69, 641-651.	6.1	36
52	Hsa-miR-196a2 polymorphism increases the risk of acute lymphoblastic leukemia in Chinese children. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014, 759, 16-21.	0.4	35
53	Expression and prognostic value of microRNA-26a and microRNA-148a in gastric cancer. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 819-827.	1.4	35
54	Body mass index (BMI) trajectories and risk of colorectal cancer in the PLCO cohort. <i>British Journal of Cancer</i> , 2018, 119, 130-132.	2.9	35

#	ARTICLE	IF	CITATIONS
55	Personal exposure to PM2.5, genetic variants and DNA damage: A multi-center population-based study in Chinese. <i>Toxicology Letters</i> , 2015, 235, 172-178.	0.4	34
56	Circulating MicroRNA-26a in Plasma and Its Potential Diagnostic Value in Gastric Cancer. <i>PLoS ONE</i> , 2016, 11, e0151345.	1.1	34
57	The prognostic significance of HOTAIR for predicting clinical outcome in patients with digestive system tumors. <i>Journal of Cancer Research and Clinical Oncology</i> , 2015, 141, 2139-2145.	1.2	33
58	Associations of IL-4, IL-4R, and IL-13 Gene Polymorphisms in Coal Workers' Pneumoconiosis in China: A Case-Control Study. <i>PLoS ONE</i> , 2011, 6, e22624.	1.1	33
59	METTL3 regulates PM2.5-induced cell injury by targeting OSGIN1 in human airway epithelial cells. <i>Journal of Hazardous Materials</i> , 2021, 415, 125573.	6.5	32
60	Cumulative effect of genome-wide association study-identified genetic variants for bladder cancer. <i>International Journal of Cancer</i> , 2014, 135, 2653-2660.	2.3	31
61	Pri-miR-34b/c rs4938723 polymorphism contributes to acute lymphoblastic leukemia susceptibility in Chinese children. <i>Leukemia and Lymphoma</i> , 2016, 57, 1436-1441.	0.6	31
62	The biogenesis and biological function of PIWI-interacting RNA in cancer. <i>Journal of Hematology and Oncology</i> , 2021, 14, 93.	6.9	31
63	Molecular epidemiology of DNA repair gene polymorphisms and head and neck cancer. <i>Journal of Biomedical Research</i> , 2013, 27, 179-92.	0.7	30
64	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018, 92, 2681-2689.	1.9	30
65	Polymorphisms of methylenetetrahydrofolate reductase and methionine synthase genes and bladder cancer risk: a case-control study with meta-analysis. <i>Clinical and Experimental Medicine</i> , 2009, 9, 9-19.	1.9	28
66	FAS and FAS Ligand Polymorphisms in the Promoter Regions and Risk of Gastric Cancer in Southern China. <i>Biochemical Genetics</i> , 2009, 47, 559-568.	0.8	28
67	A common genetic variation in the promoter of miR-107 is associated with gastric adenocarcinoma susceptibility and survival. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014, 769, 35-41.	0.4	28
68	Association between obesity and bladder cancer recurrence: A meta-analysis. <i>Clinica Chimica Acta</i> , 2018, 480, 41-46.	0.5	28
69	LncRNA <i>PCAT1</i> and its genetic variant rs1902432 are associated with prostate cancer risk. <i>Journal of Cancer</i> , 2018, 9, 1414-1420.	1.2	28
70	A functional variant in <i>TP63</i> at 3q28 associated with bladder cancer risk by creating an miR-140 binding site. <i>International Journal of Cancer</i> , 2016, 139, 65-74.	2.3	27
71	Alternative splicing related genetic variants contribute to bladder cancer risk. <i>Molecular Carcinogenesis</i> , 2020, 59, 923-929.	1.3	27
72	A Polymorphism (rs2295080) in mTOR Promoter Region and Its Association with Gastric Cancer in a Chinese Population. <i>PLoS ONE</i> , 2013, 8, e60080.	1.1	27

#	ARTICLE	IF	CITATIONS
73	Assessing the Effectiveness of Problem-Based Learning of Preventive Medicine Education in China. <i>Scientific Reports</i> , 2014, 4, 5126.	1.6	25
74	Combinations of single nucleotide polymorphisms identified in genome-wide association studies determine risk for colorectal cancer. <i>International Journal of Cancer</i> , 2019, 145, 2661-2669.	2.3	25
75	A genetic variation in the CpG island of pseudogene <i>GBAP1</i> promoter is associated with gastric cancer susceptibility. <i>Cancer</i> , 2019, 125, 2465-2473.	2.0	25
76	Polymorphism rs2682818 in miR-618 is associated with colorectal cancer susceptibility in a Han Chinese population. <i>Cancer Medicine</i> , 2018, 7, 1194-1200.	1.3	24
77	Chromosome 4p16.3 variant modify bladder cancer risk in a Chinese population. <i>Carcinogenesis</i> , 2011, 32, 872-875.	1.3	23
78	Three polymorphisms in <i>IRF6</i> and 8q24 are associated with nonsyndromic cleft lip with or without cleft palate: Evidence from 20 studies. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 3080-3086.	0.7	23
79	Association study between XPG Asp1104His polymorphism and colorectal cancer risk in a Chinese population. <i>Scientific Reports</i> , 2014, 4, 6700.	1.6	23
80	The association of rs710886 in lncRNA PCAT1 with bladder cancer risk in a Chinese population. <i>Gene</i> , 2017, 627, 226-232.	1.0	23
81	Remote modulation of lncRNA <i>GCLET</i> by risk variant at 16p13 underlying genetic susceptibility to gastric cancer. <i>Science Advances</i> , 2020, 6, eaay5525.	4.7	23
82	Effect of PM2.5 exposure on circulating fibrinogen and IL-6 levels: A systematic review and meta-analysis. <i>Chemosphere</i> , 2021, 271, 129565.	4.2	23
83	Association between circulating vitamin E and ten common cancers: evidence from large-scale Mendelian randomization analysis and a longitudinal cohort study. <i>BMC Medicine</i> , 2022, 20, 168.	2.3	23
84	The influence of genetic variants of sorafenib on clinical outcomes and toxic effects in patients with advanced renal cell carcinoma. <i>Scientific Reports</i> , 2016, 6, 20089.	1.6	22
85	Rare variants in BRCA2 and CHEK2 are associated with the risk of urinary tract cancers. <i>Scientific Reports</i> , 2016, 6, 33542.	1.6	22
86	Systematic evaluation of the effects of genetic variants on PIWI-interacting RNA expression across 33 cancer types. <i>Nucleic Acids Research</i> , 2021, 49, 90-97.	6.5	22
87	FAS rs2234767 and rs1800682 polymorphisms jointly contributed to risk of colorectal cancer by affecting SP1/STAT1 complex recruitment to chromatin. <i>Scientific Reports</i> , 2016, 6, 19229.	1.6	21
88	Fine Particulate Matter Induces Childhood Asthma Attacks via Extracellular Vesicle-Packaged Let-7i-5p-Mediated Modulation of the MAPK Signaling Pathway. <i>Advanced Science</i> , 2022, 9, e2102460.	5.6	21
89	VEGF 936C>T polymorphism and breast cancer risk: evidence from 5,729 cases and 5,868 controls. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 489-493.	1.1	20
90	The effects of particulate matters on allergic rhinitis in Nanjing, China. <i>Environmental Science and Pollution Research</i> , 2019, 26, 11452-11457.	2.7	20

#	ARTICLE	IF	CITATIONS
91	Integrative omics provide biological and clinical insights into acute respiratory distress syndrome. <i>Intensive Care Medicine</i> , 2021, 47, 761-771.	3.9	19
92	Variants in angiogenesis-related genes and the risk of clear cell renal cell carcinoma. <i>Mutagenesis</i> , 2014, 29, 419-425.	1.0	18
93	Functional POR A503V is associated with the risk of bladder cancer in a Chinese population. <i>Scientific Reports</i> , 2015, 5, 11751.	1.6	18
94	Genetic variants in N6-methyladenosine are associated with bladder cancer risk in the Chinese population. <i>Archives of Toxicology</i> , 2021, 95, 299-309.	1.9	18
95	Genetic Polymorphisms in IGF-I and IGFBP-3 Are Associated with Prostate Cancer in the Chinese Population. <i>PLoS ONE</i> , 2014, 9, e85609.	1.1	18
96	A genetic study and meta-analysis of the genetic predisposition of prostate cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 21393-21403.	0.8	18
97	Long non-coding RNA FLJ22763 is involved in the progression and prognosis of gastric cancer. <i>Gene</i> , 2019, 693, 84-91.	1.0	17
98	A genetic variant in ERCC2 is associated with gastric cancer prognosis in a Chinese population. <i>Mutagenesis</i> , 2013, 28, 441-446.	1.0	16
99	Clinical Significance of POU5F1P1 rs10505477 Polymorphism in Chinese Gastric Cancer Patients Receiving Cisplatin-Based Chemotherapy after Surgical Resection. <i>International Journal of Molecular Sciences</i> , 2014, 15, 12764-12777.	1.8	16
100	A genetic variant of miR-148a binding site in the SCRN1 3' UTR is associated with susceptibility and prognosis of gastric cancer. <i>Scientific Reports</i> , 2014, 4, 7080.	1.6	16
101	The Rare Variant rs35356162 in UHRF1BP1 Increases Bladder Cancer Risk in Han Chinese Population. <i>Frontiers in Oncology</i> , 2020, 10, 134.	1.3	16
102	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020, 137, 105543.	4.8	16
103	Metabolomics identifying biomarkers of PM2.5 exposure for vulnerable population: based on a prospective cohort study. <i>Environmental Science and Pollution Research</i> , 2021, 28, 14586-14596.	2.7	16
104	Genetic variation rs10484761 on 6p21.1 derived from a genome-wide association study is associated with gastric cancer survival in a Chinese population. <i>Gene</i> , 2014, 536, 59-64.	1.0	15
105	Hypermethylation of EIF4E promoter is associated with early onset of gastric cancer. <i>Carcinogenesis</i> , 2018, 39, 66-71.	1.3	15
106	Germline mutations in DNA repair genes are associated with bladder cancer risk and unfavourable prognosis. <i>BJU International</i> , 2018, 122, 808-813.	1.3	15
107	Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. <i>Journal of Gastroenterology</i> , 2019, 54, 939-949.	2.3	15
108	MDM2 SNP309 polymorphism is associated with colorectal cancer risk. <i>Scientific Reports</i> , 2014, 4, 4851.	1.6	14

#	ARTICLE	IF	CITATIONS
109	PSCA rs2294008 polymorphism contributes to the decreased risk for cervical cancer in a Chinese population. <i>Scientific Reports</i> , 2016, 6, 23465.	1.6	14
110	Genetic variants in PI3K/Akt/mTOR pathway genes contribute to gastric cancer risk. <i>Gene</i> , 2018, 670, 130-135.	1.0	14
111	Association study of genetic variants in estrogen metabolic pathway genes and colorectal cancer risk and survival. <i>Archives of Toxicology</i> , 2018, 92, 1991-1999.	1.9	14
112	MUC1 is associated with TFF2 methylation in gastric cancer. <i>Clinical Epigenetics</i> , 2020, 12, 37.	1.8	14
113	A prospective study of the associations among fine particulate matter, genetic variants, and the risk of colorectal cancer. <i>Environment International</i> , 2021, 147, 106309.	4.8	14
114	Genome-wide Association Study (GWAS) of Germline Copy Number Variations (CNVs) Reveal Genetic Risks of Prostate Cancer in Chinese population. <i>Journal of Cancer</i> , 2018, 9, 923-928.	1.2	13
115	Vitamin B2 intake reduces the risk for colorectal cancer: a dose-response analysis. <i>European Journal of Nutrition</i> , 2019, 58, 1591-1602.	1.8	13
116	Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. <i>Journal of Cancer</i> , 2017, 8, 1355-1361.	1.2	12
117	Evaluation of vulnerable PM2.5-exposure individuals: a repeated-measure study in an elderly population. <i>Environmental Science and Pollution Research</i> , 2018, 25, 11833-11840.	2.7	12
118	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018, 673, 174-180.	1.0	12
119	Genetic Variants in RKIP Are Associated with Clear Cell Renal Cell Carcinoma Risk in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e109285.	1.1	12
120	Functional annotation of colorectal cancer susceptibility loci identifies <i>MLH1</i> rs1800734 associated with MSI patients. <i>Gut</i> , 2016, 65, 1227-1228.	6.1	11
121	Association of Antioxidative Enzymes Polymorphisms with Efficacy of Platin and Fluorouracil-Based Adjuvant Therapy in Gastric Cancer. <i>Cellular Physiology and Biochemistry</i> , 2018, 48, 2247-2257.	1.1	11
122	Rs2262251 in lncRNA <i>RP11-462G12.2</i> is associated with nonsyndromic cleft lip with/without cleft palate. <i>Human Mutation</i> , 2019, 40, 2057-2067.	1.1	11
123	Association study between genetic variants in retinol metabolism pathway genes and prostate cancer risk. <i>Cancer Medicine</i> , 2020, 9, 9462-9470.	1.3	11
124	Genetic variations in Hippo pathway genes influence bladder cancer risk in a Chinese population. <i>Archives of Toxicology</i> , 2020, 94, 785-794.	1.9	11
125	A miR-29c binding site genetic variant in the 3'-untranslated region of LAMTOR3 gene is associated with gastric cancer risk. <i>Biomedicine and Pharmacotherapy</i> , 2015, 69, 70-75.	2.5	10
126	Identification of a novel susceptibility locus at 16q23.1 associated with childhood acute lymphoblastic leukemia in Han Chinese. <i>Human Molecular Genetics</i> , 2016, 25, ddw112.	1.4	10



#	ARTICLE	IF	CITATIONS
127	Genetic variants, PM2.5 exposure level and global DNA methylation level: A multi-center population-based study in Chinese. <i>Toxicology Letters</i> , 2017, 269, 77-82.	0.4	10
128	Tagging SNPs in the HOTAIR gene are associated with bladder cancer risk in a Chinese population. <i>Gene</i> , 2018, 664, 22-26.	1.0	10
129	Genetic Variant in Long Non-Coding RNA H19 Modulates Its Expression and Predicts Renal Cell Carcinoma Susceptibility and Mortality. <i>Frontiers in Oncology</i> , 2020, 10, 785.	1.3	10
130	Association between MLH1 -93G>A Polymorphism and Risk of Colorectal Cancer. <i>PLoS ONE</i> , 2012, 7, e50449.	1.1	9
131	A genetic variant located in the miR-532-5p-binding site of TGFBR1 is associated with the colorectal cancer risk. <i>Journal of Gastroenterology</i> , 2019, 54, 141-148.	2.3	9
132	Genetic variant in miR-621 binding sites is associated with colorectal cancer risk. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 2012-2019.	1.6	9
133	Genetic Variant rs7758229 in 6q26-q27 Is Not Associated with Colorectal Cancer Risk in a Chinese Population. <i>PLoS ONE</i> , 2013, 8, e59256.	1.1	9
134	A MAP3k1 SNP Predicts Survival of Gastric Cancer in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e96083.	1.1	9
135	Radiofrequency ablation versus partial nephrectomy for the treatment of clinical stage 1 renal masses: a systematic review and meta-analysis. <i>Chinese Medical Journal</i> , 2014, 127, 2497-503.	0.9	9
136	Associations of NR5A2 Gene Polymorphisms with the Clinicopathological Characteristics and Survival of Gastric Cancer. <i>International Journal of Molecular Sciences</i> , 2014, 15, 22902-22917.	1.8	8
137	Functional polymorphisms in apoptosis pathway genes and survival in patients with gastric cancer. <i>Environmental and Molecular Mutagenesis</i> , 2014, 55, 421-427.	0.9	8
138	Genetic variation in C12orf51 is associated with prognosis of intestinal-type gastric cancer in a Chinese population. <i>Biomedicine and Pharmacotherapy</i> , 2015, 69, 133-138.	2.5	8
139	The association analysis of <i>hOGG1</i> genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2016, 7, 66061-66068.	0.8	8
140	Identification of low-frequency variants of UGT1A3 associated with bladder cancer risk by next-generation sequencing. <i>Oncogene</i> , 2021, 40, 2382-2394.	2.6	8
141	Global internet search trends related to gastrointestinal symptoms predict regional COVID-19 outbreaks. <i>Journal of Infection</i> , 2022, 84, 56-63.	1.7	8
142	Evaluation of genome-wide genotyping concordance between tumor tissues and peripheral blood. <i>Genomics</i> , 2017, 109, 108-112.	1.3	7
143	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018, 33, 82-87.	2.7	7
144	Genetic variants in SMARC genes are associated with DNA damage levels in Chinese population. <i>Toxicology Letters</i> , 2014, 229, 327-332.	0.4	6

#	ARTICLE	IF	CITATIONS
145	Genetic variation in IGF1 predicts renal cell carcinoma susceptibility and prognosis in Chinese population. <i>Scientific Reports</i> , 2016, 6, 39014.	1.6	6
146	Genetic variants in multisynthetase complex genes are associated with DNA damage levels in Chinese populations. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2016, 786, 8-13.	0.4	6
147	Information transduction capacity reduces the uncertainties in annotation-free isoform discovery and quantification. <i>Nucleic Acids Research</i> , 2017, 45, e143-e143.	6.5	6
148	Probabilistic natural mapping of gene-level tests for genome-wide association studies. <i>Briefings in Bioinformatics</i> , 2018, 19, 545-553.	3.2	6
149	Explaining the Genetic Causality for Complex Phenotype via Deep Association Kernel Learning. <i>Patterns</i> , 2020, 1, 100057.	3.1	6
150	Association of genetic variants in autophagy-lysosome pathway genes with susceptibility and survival to prostate cancer. <i>Gene</i> , 2022, 808, 145953.	1.0	6
151	Long-term risk of colorectal cancer after removal of adenomas during screening colonoscopies in a large community-based population in China. <i>International Journal of Cancer</i> , 2022, 150, 594-602.	2.3	6
152	Effects of TSP-1 -696 C/T polymorphism on bladder cancer susceptibility and clinicopathologic features. <i>Cancer Genetics</i> , 2014, 207, 247-252.	0.2	5
153	Identification of novel susceptibility loci for non-syndromic cleft lip with or without cleft palate. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 13669-13678.	1.6	5
154	Novel CpG-SNPs in the gastric acid secretion pathway GNAI3 and susceptibility to gastric cancer. <i>Gene</i> , 2020, 736, 144447.	1.0	5
155	Genetic variations in the CTLA-4 immune checkpoint pathway are associated with colon cancer risk, prognosis, and immune infiltration via regulation of IQCB1 expression. <i>Archives of Toxicology</i> , 2021, 95, 2053-2063.	1.9	5
156	CoSMED: a user-friendly web server to estimate 5-year survival probability of left-sided and right-sided colorectal cancer patients using molecular data. <i>Bioinformatics</i> , 2021, 38, 278-281.	1.8	5
157	Genetic Variations in the 3'-untranslated Regions of Genes Involved in the Cell Cycle and Apoptosis Pathways Affect Bladder Cancer Risk. <i>Cancer Genomics and Proteomics</i> , 2018, 15, 67-72.	1.0	5
158	Genetic variants in Ras/Raf/MEK/ERK pathway are associated with gastric cancer risk in Chinese Han population. <i>Archives of Toxicology</i> , 2020, 94, 2683-2690.	1.9	4
159	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021, 95, 2523-2532.	1.9	4
160	Association Between MIF-AS rs755622 and Nephrolithiasis Risk in a Chinese Population. <i>Medical Science Monitor</i> , 2016, 22, 563-568.	0.5	4
161	High-density lipoprotein, low-density lipoprotein and triglyceride levels and upper gastrointestinal cancers risk: a trans-ancestry Mendelian randomization study. <i>European Journal of Clinical Nutrition</i> , 2022, , .	1.3	4
162	TSP-1-1223 A/G Polymorphism as a Potential Predictor of the Recurrence Risk of Bladder Cancer in a Chinese Population. <i>International Journal of Genomics</i> , 2013, 2013, 1-9.	0.8	3

#	ARTICLE	IF	CITATIONS
163	Genetic variants of H2AX gene were associated with P M 2.5 -modulated DNA damage levels in Chinese Han populations. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2015, 778, 41-45.	0.4	3
164	Validation of the novel susceptibility loci for prostate cancer in a Chinese population. Oncology Letters, 2017, 15, 2567-2573.	0.8	3
165	Genetic variants in XDH are associated with prognosis for gastric cancer in a Chinese population. Gene, 2018, 663, 196-202.	1.0	3
166	Polymorphism rs4787951 in IL-4R contributes to the increased risk of renal cell carcinoma in a Chinese population. Gene, 2019, 685, 242-247.	1.0	3
167	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. Archives of Toxicology, 2020, 94, 325-333.	1.9	3
168	A transcriptomic study for identifying cardia-specific and non-specific gastric cancer prognostic factors using genetic algorithm-based methods. Journal of Cellular and Molecular Medicine, 2020, 24, 9457-9465.	1.6	3
169	Genetic variants in Hippo signalling pathway-related genes affect the risk of colorectal cancer. Archives of Toxicology, 2021, 95, 271-281.	1.9	3
170	Identification of common genetic variants associated with serum concentrations of p, p <sup>2</sup> -DDE in non-occupational populations in eastern China. Environment International, 2021, 152, 106507.	4.8	3
171	Genome-Wide Association Analyses Identify <i>CATSPER1</i> as a Mediator of Colorectal Cancer Susceptibility and Progression. Cancer Research, 2022, 82, 986-997.	0.4	3
172	Evaluation of genetic variants in nucleosome remodeling and deacetylase (NuRD) complex subunits encoding genes and gastric cancer susceptibility. Archives of Toxicology, 2022, 96, 1739-1749.	1.9	2
173	Cohort profile: The National Colorectal Cancer Cohort (NCRCC) study in China. BMJ Open, 2021, 11, e051397.	0.8	2
174	Genetic variants in the Folic acid Metabolic Pathway Genes predict outcomes of metastatic Colorectal Cancer patients receiving first-line Chemotherapy. Journal of Cancer, 2020, 11, 6507-6515.	1.2	1
175	Genetic variants in splicing factor genes and susceptibility to bladder cancer. Gene, 2022, 809, 146022.	1.0	1
176	SOD2 rs4880 CT/CC genotype to predict poor survival for Chinese gastric cancer patients received platinum and fluorouracil based adjuvant chemotherapy.. Journal of Clinical Oncology, 2015, 33, 11037-11037.	0.8	1
177	Comprehensive genetic mutation analysis of human gastric adenocarcinomas.. Journal of Clinical Oncology, 2013, 31, 4106-4106.	0.8	0
178	Single nucleotide polymorphism of SOD2 to predict survival for Chinese gastric cancer patients received platinum/fluorouracil-based adjuvant chemotherapy.. Journal of Clinical Oncology, 2014, 32, e15035-e15035.	0.8	0
179	Genetic variants in the Hedgehog signaling pathway genes are associated with gastric cancer risk in a Chinese Han population. Journal of Biomedical Research, 2022, 36, 22.	0.7	0