Meilin Wang

List of Publications by Citations

176	4,127	36	56
papers	citations	h-index	g-index
182	4,983	7.3	5.24
ext. papers	ext. citations	avg, IF	L-index

#	Paper	IF	Citations
176	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. *Nature Genetics*, **2011**, 43, 1215-8	36.3	215
175	LncRNA MT1JP functions as a ceRNA in regulating FBXW7 through competitively binding to miR-92a-3p in gastric cancer. *Molecular Cancer*, **2018**, 17, 87	42.1	166
174	Exosome-transmitted long non-coding RNA PTENP1 suppresses bladder cancer progression. *Molecular Cancer*, **2018**, 17, 143	42.1	134
173	Genome-wide association study identifies a new susceptibility locus for cleft lip with or without a cleft palate. *Nature Communications*, **2015**, 6, 6414	17.4	124
172	Genetic variants in lncRNA HOTAIR are associated with risk of colorectal cancer. *Mutagenesis*, **2015**, 30, 303-10	2.8	112
171	A functional polymorphism in MSMB gene promoter is associated with prostate cancer risk and serum MSMB expression. *Prostate*, **2010**, 70, 1146-52	4.2	98
170	Identification of novel piRNAs in bladder cancer. *Cancer Letters*, **2015**, 356, 561-7	9.9	91
169	The association analysis of lncRNA HOTAIR genetic variants and gastric cancer risk in a Chinese population. *Oncotarget*, **2015**, 6, 31255-62	3.3	91
168	Circulating miR-497 and miR-663b in plasma are potential novel biomarkers for bladder cancer. *Scientific Reports*, **2015**, 5, 10437	4.9	87
167	Multiomics Evaluation of Gastrointestinal and Other Clinical Characteristics of COVID-19. *Gastroenterology*, **2020**, 158, 2298-2301.e7	13.3	83
166	Polymorphism of the pre-miR-146a is associated with risk of cervical cancer in a Chinese population. *Gynecologic Oncology*, **2011**, 122, 33-7	4.9	82
165	Association of genetic variants in lncRNA H19 with risk of colorectal cancer in a Chinese population. *Oncotarget*, **2016**, 7, 25470-7	3.3	68
164	Genetic variants in miRNAs predict bladder cancer risk and recurrence. *Cancer Research*, **2012**, 72, 6173-82	10.1	67
163	Genome-wide association analysis of Vogt-Koyanagi-Harada syndrome identifies two new susceptibility loci at 1p31.2 and 10q21.3. *Nature Genetics*, **2014**, 46, 1007-11	36.3	65
162	Circular RNAs in body fluids as cancer biomarkers: the new frontier of liquid biopsies. *Molecular Cancer*, **2021**, 20, 13	42.1	64
161	Genetic variants in lncRNA H19 are associated with the risk of bladder cancer in a Chinese population. *Mutagenesis*, **2016**, 31, 531-8	2.8	60
160	A functional polymorphism in miRNA-196a2 is associated with colorectal cancer risk in a Chinese population. *DNA and Cell Biology*, **2012**, 31, 350-4	3.6	56

159	A functional polymorphism in Pre-miR-146a is associated with susceptibility to gastric cancer in a Chinese population. *DNA and Cell Biology*, **2012**, 31, 1290-5	3.6	56
158	A genetic variant in miR-146a modifies colorectal cancer susceptibility in a Chinese population. *Archives of Toxicology*, **2013**, 87, 825-33	5.8	55
157	Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. *Gene*, **2015**, 556, 227-34	3.8	53
156	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. *Gut*, **2017**, 66, 581-587	19.2	51
155	Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. *Carcinogenesis*, **2014**, 35, 2723-30	4.6	51
154	A novel antisense long noncoding RNA regulates the expression of MDC1 in bladder cancer. *Oncotarget*, **2015**, 6, 484-93	3.3	50
153	Genetic variant in PSCA predicts survival of diffuse-type gastric cancer in a Chinese population. *International Journal of Cancer*, **2011**, 129, 1207-13	7.5	49
152	Global gene expression profiling of human bronchial epithelial cells exposed to airborne fine particulate matter collected from Wuhan, China. *Toxicology Letters*, **2014**, 228, 25-33	4.4	48
151	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. *Nature Communications*, **2016**, 7, 11478	17.4	45
150	The classic EDCs, phthalate esters and organochlorines, in relation to abnormal sperm quality: a systematic review with meta-analysis. *Scientific Reports*, **2016**, 6, 19982	4.9	44
149	Genetic variants in noncoding PIWI-interacting RNA and colorectal cancer risk. *Cancer*, **2015**, 121, 2044-52	6.4	43
148	Genetic variations in microRNAs and the risk and survival of renal cell cancer. *Carcinogenesis*, **2014**, 35, 1629-35	4.6	41
147	miR-107 regulates tumor progression by targeting NF1 in gastric cancer. *Scientific Reports*, **2016**, 6, 36531	4.9	40
146	Clinical significance of SOD2 and GSTP1 gene polymorphisms in Chinese patients with gastric cancer. *Cancer*, **2012**, 118, 5489-96	6.4	40
145	Folic acid supplements and colorectal cancer risk: meta-analysis of randomized controlled trials. *Scientific Reports*, **2015**, 5, 12044	4.9	39
144	Association of three polymorphisms in ARID5B, IKZF1 and CEBPE with the risk of childhood acute lymphoblastic leukemia in a Chinese population. *Gene*, **2013**, 524, 203-7	3.8	38
143	A novel functional polymorphism C1797G in the MDM2 promoter is associated with risk of bladder cancer in a Chinese population. *Clinical Cancer Research*, **2008**, 14, 3633-40	12.9	38
142	Large-scale association analysis in Asians identifies new susceptibility loci for prostate cancer. *Nature Communications*, **2015**, 6, 8469	17.4	37

141	Common genetic variants in pre-microRNAs are associated with risk of coal workersS pneumoconiosis. *Journal of Human Genetics*, **2010**, 55, 13-7	4.3	37
140	Meta-analysis on the effectiveness of team-based learning on medical education in China. *BMC Medical Education*, **2018**, 18, 77	3.3	35
139	Replication and cumulative effects of GWAS-identified genetic variations for prostate cancer in Asians: a case-control study in the ChinaPCa consortium. *Carcinogenesis*, **2012**, 33, 356-60	4.6	33
138	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. *Gastroenterology*, **2017**, 152, 2011-2021	13.3	32
137	Genetic variants in m6A modification genes are associated with colorectal cancer risk. *Carcinogenesis*, **2020**, 41, 8-17	4.6	30
136	A functional variant in miR-143 promoter contributes to prostate cancer risk. *Archives of Toxicology*, **2016**, 90, 403-14	5.8	30
135	An inverse association between tea consumption and colorectal cancer risk. *Oncotarget*, **2017**, 8, 37367-37376	3.3	30
134	Circulating MicroRNA-26a in Plasma and Its Potential Diagnostic Value in Gastric Cancer. *PLoS ONE*, **2016**, 11, e0151345	3.7	30
133	Expression and prognostic value of microRNA-26a and microRNA-148a in gastric cancer. *Journal of Gastroenterology and Hepatology (Australia)*, **2017**, 32, 819-827	4	29
132	The prognostic significance of HOTAIR for predicting clinical outcome in patients with digestive system tumors. *Journal of Cancer Research and Clinical Oncology*, **2015**, 141, 2139-45	4.9	29
131	The HOTAIR, PRNCR1 and POLR2E polymorphisms are associated with cancer risk: a meta-analysis. *Oncotarget*, **2017**, 8, 43271-43283	3.3	29
130	Genome-Wide Association Study of Bladder Cancer in a Chinese Cohort Reveals a New Susceptibility Locus at 5q12.3. *Cancer Research*, **2016**, 76, 3277-84	10.1	29
129	Environmental factors, seven GWAS-identified susceptibility loci, and risk of gastric cancer and its precursors in a Chinese population. *Cancer Medicine*, **2017**, 6, 708-720	4.8	28
128	Hsa-miR-196a2 polymorphism increases the risk of acute lymphoblastic leukemia in Chinese children. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2014**, 759, 16-21	3.3	28
127	Personal exposure to PM2.5, genetic variants and DNA damage: a multi-center population-based study in Chinese. *Toxicology Letters*, **2015**, 235, 172-8	4.4	26
126	FAS and FAS ligand polymorphisms in the promoter regions and risk of gastric cancer in Southern China. *Biochemical Genetics*, **2009**, 47, 559-68	2.4	26
125	Associations of IL-4, IL-4R, and IL-13 gene polymorphisms in coal workersSpneumoconiosis in China: a case-control study. *PLoS ONE*, **2011**, 6, e22624	3.7	26
124	A common genetic variation in the promoter of miR-107 is associated with gastric adenocarcinoma susceptibility and survival. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2014**, 769, 35-41	3.3	25

123	KCNMA1 cooperating with PTK2 is a novel tumor suppressor in gastric cancer and is associated with disease outcome. *Molecular Cancer*, **2017**, 16, 46	42.1	24
122	Cumulative effect of genome-wide association study-identified genetic variants for bladder cancer. *International Journal of Cancer*, **2014**, 135, 2653-60	7.5	24
121	Short-term effects of ambient air pollution and childhood lower respiratory diseases. *Scientific Reports*, **2017**, 7, 4414	4.9	24
120	Molecular epidemiology of DNA repair gene polymorphisms and head and neck cancer. *Journal of Biomedical Research*, **2013**, 27, 179-92	1.5	24
119	A polymorphism (rs2295080) in mTOR promoter region and its association with gastric cancer in a Chinese population. *PLoS ONE*, **2013**, 8, e60080	3.7	24
118	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. *Archives of Toxicology*, **2018**, 92, 2681-2689	5.8	24
117	Pri-miR-34b/c rs4938723 polymorphism contributes to acute lymphoblastic leukemia susceptibility in Chinese children. *Leukemia and Lymphoma*, **2016**, 57, 1436-41	1.9	23
116	The association of rs710886 in lncRNA PCAT1 with bladder cancer risk in a Chinese population. *Gene*, **2017**, 627, 226-232	3.8	22
115	LncRNA and its genetic variant rs1902432 are associated with prostate cancer risk. *Journal of Cancer*, **2018**, 9, 1414-1420	4.5	22
114	Three polymorphisms in IRF6 and 8q24 are associated with nonsyndromic cleft lip with or without cleft palate: evidence from 20 studies. *American Journal of Medical Genetics, Part A*, **2012**, 158A, 3080-6	2.5	22
113	Environmental exposure to BDE47 is associated with increased diabetes prevalence: Evidence from community-based case-control studies and an animal experiment. *Scientific Reports*, **2016**, 6, 27854	4.9	22
112	Polymorphisms of methylenetetrahydrofolate reductase and methionine synthase genes and bladder cancer risk: a case-control study with meta-analysis. *Clinical and Experimental Medicine*, **2009**, 9, 9-19	4.9	21
111	Association study between XPG Asp1104His polymorphism and colorectal cancer risk in a Chinese population. *Scientific Reports*, **2014**, 4, 6700	4.9	20
110	VEGF 936C>T polymorphism and breast cancer risk: evidence from 5,729 cases and 5,868 controls. *Breast Cancer Research and Treatment*, **2011**, 125, 489-93	4.4	20
109	Chromosome 4p16.3 variant modify bladder cancer risk in a Chinese population. *Carcinogenesis*, **2011**, 32, 872-5	4.6	20
108	FAS rs2234767 and rs1800682 polymorphisms jointly contributed to risk of colorectal cancer by affecting SP1/STAT1 complex recruitment to chromatin. *Scientific Reports*, **2016**, 6, 19229	4.9	19
107	A functional variant in TP63 at 3q28 associated with bladder cancer risk by creating an miR-140-5p binding site. *International Journal of Cancer*, **2016**, 139, 65-74	7.5	19
106	Genome-wide long non-coding RNAs identified a panel of novel plasma biomarkers for gastric cancer diagnosis. *Gastric Cancer*, **2019**, 22, 731-741	7.6	19

105	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. *Gut*, **2020**, 69, 641-651	19.2	18
104	A genetic study and meta-analysis of the genetic predisposition of prostate cancer in a Chinese population. *Oncotarget*, **2016**, 7, 21393-403	3.3	17
103	Association between obesity and bladder cancer recurrence: A meta-analysis. *Clinica Chimica Acta*, **2018**, 480, 41-46	6.2	16
102	Rare variants in BRCA2 and CHEK2 are associated with the risk of urinary tract cancers. *Scientific Reports*, **2016**, 6, 33542	4.9	16
101	A genetic variant of miR-148a binding site in the SCRN1 3SUTR is associated with susceptibility and prognosis of gastric cancer. *Scientific Reports*, **2014**, 4, 7080	4.9	15
100	Body mass index (BMI) trajectories and risk of colorectal cancer in the PLCO cohort. *British Journal of Cancer*, **2018**, 119, 130-132	8.7	15
99	Clinical significance of POU5F1P1 rs10505477 polymorphism in Chinese gastric cancer patients receving cisplatin-based chemotherapy after surgical resection. *International Journal of Molecular Sciences*, **2014**, 15, 12764-77	6.3	15
98	A genetic variant in ERCC2 is associated with gastric cancer prognosis in a Chinese population. *Mutagenesis*, **2013**, 28, 441-6	2.8	15
97	Genetic polymorphisms in IGF-I and IGFBP-3 are associated with prostate cancer in the Chinese population. *PLoS ONE*, **2014**, 9, e85609	3.7	15
96	The influence of genetic variants of sorafenib on clinical outcomes and toxic effects in patients with advanced renal cell carcinoma. *Scientific Reports*, **2016**, 6, 20089	4.9	14
95	Genetic variation rs10484761 on 6p21.1 derived from a genome-wide association study is associated with gastric cancer survival in a Chinese population. *Gene*, **2014**, 536, 59-64	3.8	14
94	Polymorphism rs2682818 in miR-618 is associated with colorectal cancer susceptibility in a Han Chinese population. *Cancer Medicine*, **2018**, 7, 1194-1200	4.8	13
93	PSCA rs2294008 polymorphism contributes to the decreased risk for cervical cancer in a Chinese population. *Scientific Reports*, **2016**, 6, 23465	4.9	13
92	Functional POR A503V is associated with the risk of bladder cancer in a Chinese population. *Scientific Reports*, **2015**, 5, 11751	4.9	13
91	Assessing the effectiveness of problem-based learning of preventive medicine education in China. *Scientific Reports*, **2014**, 4, 5126	4.9	13
90	Variants in angiogenesis-related genes and the risk of clear cell renal cell carcinoma. *Mutagenesis*, **2014**, 29, 419-25	2.8	13
89	Combinations of single nucleotide polymorphisms identified in genome-wide association studies determine risk for colorectal cancer. *International Journal of Cancer*, **2019**, 145, 2661-2669	7.5	12
88	MDM2 SNP309 polymorphism is associated with colorectal cancer risk. *Scientific Reports*, **2014**, 4, 4851	4.9	12

87	Hypermethylation of EIF4E promoter is associated with early onset of gastric cancer. *Carcinogenesis* , **2018**, 39, 66-71	4.6	12
86	Germline mutations in DNA repair genes are associated with bladder cancer risk and unfavourable prognosis. *BJU International*, **2018**, 122, 808-813	5.6	12
85	Genetic variants in PI3K/Akt/mTOR pathway genes contribute to gastric cancer risk. *Gene*, **2018**, 670, 130-135	3.8	12
84	Association study of genetic variants in estrogen metabolic pathway genes and colorectal cancer risk and survival. *Archives of Toxicology*, **2018**, 92, 1991-1999	5.8	12
83	A genetic variation in the CpG island of pseudogene GBAP1 promoter is associated with gastric cancer susceptibility. *Cancer*, **2019**, 125, 2465-2473	6.4	11
82	The effects of particulate matters on allergic rhinitis in Nanjing, China. *Environmental Science and Pollution Research*, **2019**, 26, 11452-11457	5.1	11
81	Functional annotation of colorectal cancer susceptibility loci identifies MLH1 rs1800734 associated with MSI patients. *Gut*, **2016**, 65, 1227-8	19.2	11
80	Effect of PM exposure on circulating fibrinogen and IL-6 levels: A systematic review and meta-analysis. *Chemosphere*, **2021**, 271, 129565	8.4	11
79	Genetic variants, PM exposure level and global DNA methylation level: A multi-center population-based study in Chinese. *Toxicology Letters*, **2017**, 269, 77-82	4.4	10
78	Remote modulation of lncRNA by risk variant at 16p13 underlying genetic susceptibility to gastric cancer. *Science Advances*, **2020**, 6, eaay5525	14.3	10
77	Alternative splicing related genetic variants contribute to bladder cancer risk. *Molecular Carcinogenesis*, **2020**, 59, 923-929	5	10
76	Genome-wide Association Study (GWAS) of Germline Copy Number Variations (CNVs) Reveal Genetic Risks of Prostate Cancer in Chinese population. *Journal of Cancer*, **2018**, 9, 923-928	4.5	10
75	Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. *Journal of Cancer*, **2017**, 8, 1355-1361	4.5	10
74	Genetic variants in RKIP are associated with clear cell renal cell carcinoma risk in a Chinese population. *PLoS ONE*, **2014**, 9, e109285	3.7	10
73	A miR-29c binding site genetic variant in the 3Suntranslated region of LAMTOR3 gene is associated with gastric cancer risk. *Biomedicine and Pharmacotherapy*, **2015**, 69, 70-5	7.5	9
72	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. *Gene*, **2018**, 673, 174-180	3.8	9
71	A genetic variant located in the miR-532-5p-binding site of TGFBR1 is associated with the colorectal cancer risk. *Journal of Gastroenterology*, **2019**, 54, 141-148	6.9	9
70	Association between MLH1 -93G>a polymorphism and risk of colorectal cancer. *PLoS ONE*, **2012**, 7, e50449	3.7	9

69	Genetic variant rs7758229 in 6q26-q27 is not associated with colorectal cancer risk in a Chinese population. *PLoS ONE*, **2013**, 8, e59256	3.7	9
68	Long non-coding RNA FLJ22763 is involved in the progression and prognosis of gastric cancer. *Gene* , **2019**, 693, 84-91	3.8	8
67	Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. *Journal of Gastroenterology*, **2019**, 54, 939-949	6.9	8
66	Genetic variations in Hippo pathway genes influence bladder cancer risk in a Chinese population. *Archives of Toxicology*, **2020**, 94, 785-794	5.8	8
65	Tagging SNPs in the HOTAIR gene are associated with bladder cancer risk in a Chinese population. *Gene*, **2018**, 664, 22-26	3.8	8
64	Functional polymorphisms in apoptosis pathway genes and survival in patients with gastric cancer. *Environmental and Molecular Mutagenesis*, **2014**, 55, 421-7	3.2	8
63	Associations of NR5A2 gene polymorphisms with the clinicopathological characteristics and survival of gastric cancer. *International Journal of Molecular Sciences*, **2014**, 15, 22902-17	6.3	8
62	Systematic evaluation of the effects of genetic variants on PIWI-interacting RNA expression across 33 cancer types. *Nucleic Acids Research*, **2021**, 49, 90-97	20.1	8
61	Radiofrequency ablation versus partial nephrectomy for the treatment of clinical stage 1 renal masses: a systematic review and meta-analysis. *Chinese Medical Journal*, **2014**, 127, 2497-503	2.9	8
60	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. *Environment International*, **2020**, 137, 105543	12.9	7
59	Genetic variation in C12orf51 is associated with prognosis of intestinal-type gastric cancer in a Chinese population. *Biomedicine and Pharmacotherapy*, **2015**, 69, 133-8	7.5	7
58	Association study between genetic variants in retinol metabolism pathway genes and prostate cancer risk. *Cancer Medicine*, **2020**, 9, 9462-9470	4.8	7
57	The association analysis of hOGG1 genetic variants and gastric cancer risk in a Chinese population. *Oncotarget*, **2016**, 7, 66061-66068	3.3	7
56	METTL3 regulates PM-induced cell injury by targeting OSGIN1 in human airway epithelial cells. *Journal of Hazardous Materials*, **2021**, 415, 125573	12.8	7
55	Evaluation of genome-wide genotyping concordance between tumor tissues and peripheral blood. *Genomics*, **2017**, 109, 108-112	4.3	6
54	Evaluation of vulnerable PM-exposure individuals: a repeated-measure study in an elderly population. *Environmental Science and Pollution Research*, **2018**, 25, 11833-11840	5.1	6
53	Genetic variants in multisynthetase complex genes are associated with DNA damage levels in Chinese populations. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2016**, 786, 8-13	3.3	6
52	Association of Antioxidative Enzymes Polymorphisms with Efficacy of Platin and Fluorouracil-Based Adjuvant Therapy in Gastric Cancer. *Cellular Physiology and Biochemistry*, **2018**, 48, 2247-2257	3.9	6

51	Information transduction capacity reduces the uncertainties in annotation-free isoform discovery and quantification. *Nucleic Acids Research*, **2017**, 45, e143	20.1	6
50	A MAP3k1 SNP predicts survival of gastric cancer in a Chinese population. *PLoS ONE*, **2014**, 9, e96083	3.7	6
49	Identification of a novel susceptibility locus at 16q23.1 associated with childhood acute lymphoblastic leukemia in Han Chinese. *Human Molecular Genetics*, **2016**, 25, 2873-2880	5.6	6
48	Genetic variant in miR-21 binding sites is associated with colorectal cancer risk. *Journal of Cellular and Molecular Medicine*, **2019**, 23, 2012-2019	5.6	6
47	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction.. *Molecular Cancer*, **2022**, 21, 49	42.1	6
46	The Rare Variant rs35356162 in Increases Bladder Cancer Risk in Han Chinese Population. *Frontiers in Oncology*, **2020**, 10, 134	5.3	5
45	Probabilistic natural mapping of gene-level tests for genome-wide association studies. *Briefings in Bioinformatics*, **2018**, 19, 545-553	13.4	5
44	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. *EBioMedicine*, **2018**, 33, 82-87	8.8	5
43	Vitamin B intake reduces the risk for colorectal cancer: a dose-response analysis. *European Journal of Nutrition*, **2019**, 58, 1591-1602	5.2	5
42	Rs2262251 in lncRNA RP11-462G12.2 is associated with nonsyndromic cleft lip with/without cleft palate. *Human Mutation*, **2019**, 40, 2057-2067	4.7	5
41	Genetic variants in SMARC genes are associated with DNA damage levels in Chinese population. *Toxicology Letters*, **2014**, 229, 327-32	4.4	5
40	Effects of TSP-1-696 C/T polymorphism on bladder cancer susceptibility and clinicopathologic features. *Cancer Genetics*, **2014**, 207, 247-52	2.3	5
39	Genetic Variations in the 3Suntranslated Regions of Genes Involved in the Cell Cycle and Apoptosis Pathways Affect Bladder Cancer Risk. *Cancer Genomics and Proteomics*, **2018**, 15, 67-72	3.3	5
38	Genetic variation in IGF1 predicts renal cell carcinoma susceptibility and prognosis in Chinese population. *Scientific Reports*, **2016**, 6, 39014	4.9	5
37	Metabolomics identifying biomarkers of PM exposure for vulnerable population: based on a prospective cohort study. *Environmental Science and Pollution Research*, **2021**, 28, 14586-14596	5.1	5
36	Genetic variants in N6-methyladenosine are associated with bladder cancer risk in the Chinese population. *Archives of Toxicology*, **2021**, 95, 299-309	5.8	5
35	A prospective study of the associations among fine particulate matter, genetic variants, and the risk of colorectal cancer. *Environment International*, **2021**, 147, 106309	12.9	4
34	Genetic variants of H2AX gene were associated with PM2.5-modulated DNA damage levels in Chinese Han populations. *Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis*, **2015**, 778, 41-5	3.3	3

33	Genetic Variant in Long Non-Coding RNA Modulates Its Expression and Predicts Renal Cell Carcinoma Susceptibility and Mortality. *Frontiers in Oncology*, **2020**, 10, 785	5.3	3
32	MUC1 is associated with TFF2 methylation in gastric cancer. *Clinical Epigenetics*, **2020**, 12, 37	7.7	3
31	Novel CpG-SNPs in the gastric acid secretion pathway GNAI3 and susceptibility to gastric cancer. *Gene*, **2020**, 736, 144447	3.8	3
30	Validation of the novel susceptibility loci for prostate cancer in a Chinese population. *Oncology Letters*, **2018**, 15, 2567-2573	2.6	3
29	TSP-1-1223 A/G Polymorphism as a Potential Predictor of the Recurrence Risk of Bladder Cancer in a Chinese Population. *International Journal of Genomics*, **2013**, 2013, 473242	2.5	3
28	Explaining the Genetic Causality for Complex Phenotype via Deep Association Kernel Learning. *Patterns*, **2020**, 1, 100057	5.1	3
27	The biogenesis and biological function of PIWI-interacting RNA in cancer. *Journal of Hematology and Oncology*, **2021**, 14, 93	22.4	3
26	Identification of low-frequency variants of UGT1A3 associated with bladder cancer risk by next-generation sequencing. *Oncogene*, **2021**, 40, 2382-2394	9.2	3
25	Genetic variants in Ras/Raf/MEK/ERK pathway are associated with gastric cancer risk in Chinese Han population. *Archives of Toxicology*, **2020**, 94, 2683-2690	5.8	2
24	A transcriptomic study for identifying cardia- and non-cardia-specific gastric cancer prognostic factors using genetic algorithm-based methods. *Journal of Cellular and Molecular Medicine*, **2020**, 24, 9457-9465	5.6	2
23	Association Between MIF-AS rs755622 and Nephrolithiasis Risk in a Chinese Population. *Medical Science Monitor*, **2016**, 22, 563-8	3.2	2
22	Fine Particulate Matter Induces Childhood Asthma Attacks via Extracellular Vesicle-Packaged Let-7i-5p-Mediated Modulation of the MAPK Signaling Pathway. *Advanced Science*, **2021**, e2102460	13.6	2
21	Global internet search trends related to gastrointestinal symptoms predict regional COVID-19 outbreaks. *Journal of Infection*, **2021**,	18.9	2
20	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. *Archives of Toxicology*, **2020**, 94, 325-333	5.8	2
19	Integrative omics provide biological and clinical insights into acute respiratory distress syndrome. *Intensive Care Medicine*, **2021**, 47, 761-771	14.5	2
18	Polymorphism rs4787951 in IL-4R contributes to the increased risk of renal cell carcinoma in a Chinese population. *Gene*, **2019**, 685, 242-247	3.8	2
17	Genetic variants in Hippo signalling pathway-related genes affect the risk of colorectal cancer. *Archives of Toxicology*, **2021**, 95, 271-281	5.8	2
16	Association of genetic variants in autophagy-lysosome pathway genes with susceptibility and survival to prostate cancer. *Gene*, **2022**, 808, 145953	3.8	2

List of Publications

15	Association between circulating vitamin E and ten common cancers: evidence from large-scale Mendelian randomization analysis and a longitudinal cohort study.. *BMC Medicine*, **2022**, 20, 168	11.4	2
14	Genetic variants in XDH are associated with prognosis for gastric cancer in a Chinese population. *Gene*, **2018**, 663, 196-202	3.8	1
13	Identification of novel susceptibility loci for non-syndromic cleft lip with or without cleft palate. *Journal of Cellular and Molecular Medicine*, **2020**, 24, 13669-13678	5.6	1
12	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. *Archives of Toxicology*, **2021**, 95, 2523-2532	5.8	1
11	Long-term risk of colorectal cancer after removal of adenomas during screening colonoscopies in a large community-based population in China. *International Journal of Cancer*, **2021**, 150, 594	7.5	0
10	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	2.2	0
9	Genetic variations in the CTLA-4 immune checkpoint pathway are associated with colon cancer risk, prognosis, and immune infiltration via regulation of IQCB1 expression. *Archives of Toxicology*, **2021**, 95, 2053-2063	5.8	0
8	Identification of common genetic variants associated with serum concentrations of p, pSDDE in non-occupational populations in eastern China. *Environment International*, **2021**, 152, 106507	12.9	0
7	Evaluation of genetic variants in nucleosome remodeling and deacetylase (NuRD) complex subunits encoding genes and gastric cancer susceptibility.. *Archives of Toxicology*, **2022**, 1	5.8	0
6	Cohort profile: The National Colorectal Cancer Cohort (NCRCC) study in China.. *BMJ Open*, **2021**, 11, e051397	3	0
5	Genetic variants in the Hedgehog signaling pathway genes are associated with gastric cancer risk in a Chinese Han population.. *Journal of Biomedical Research*, **2021**, 36, 22-31	1.5	
4	Genetic variants in splicing factor genes and susceptibility to bladder cancer. *Gene*, **2022**, 809, 146022	3.8	
3	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	2.2	
2	Comprehensive genetic mutation analysis of human gastric adenocarcinomas.. *Journal of Clinical Oncology*, **2013**, 31, 4106-4106	2.2	
1	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	4.5	