

Zhengdong Zhang

List of Publications by Year in descending order

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158
papers

4,462
citations

117625

34
h-index

149698

56
g-index

160
all docs

160
docs citations

160
times ranked

6851
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.	21.4	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.	19.2	218
3	Exosome-transmitted long non-coding RNA PTENP1 suppresses bladder cancer progression. <i>Molecular Cancer</i> , 2018, 17, 143.	19.2	217
4	Genetic variants in lncRNA HOTAIR are associated with risk of colorectal cancer. <i>Mutagenesis</i> , 2015, 30, 303-310.	2.6	128
5	Multiomics Evaluation of Gastrointestinal and Other Clinical Characteristics of COVID-19. <i>Gastroenterology</i> , 2020, 158, 2298-2301.e7.	1.3	117
6	Identification of novel piRNAs in bladder cancer. <i>Cancer Letters</i> , 2015, 356, 561-567.	7.2	115
7	Circulating miR-497 and miR-663b in plasma are potential novel biomarkers for bladder cancer. <i>Scientific Reports</i> , 2015, 5, 10437.	3.3	105
8	The association analysis of lncRNA HOTAIR genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2015, 6, 31255-31262.	1.8	95
9	Association of genetic variants in lncRNA H19 with risk of colorectal cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 25470-25477.	1.8	90
10	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016, 7, 11478.	12.8	73
11	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction. <i>Molecular Cancer</i> , 2022, 21, 49.	19.2	72
12	Genetic variants in lncRNA H19 are associated with the risk of bladder cancer in a Chinese population. <i>Mutagenesis</i> , 2016, 31, 531-538.	2.6	70
13	Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. <i>Gene</i> , 2015, 556, 227-234.	2.2	66
14	Polymorphisms of Methionine Synthase and Methionine Synthase Reductase and Risk of Squamous Cell Carcinoma of the Head and Neck: a Case-Control Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 1188-1193.	2.5	61
15	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.	0.8	58
16	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017, 152, 2011-2021.	1.3	58
17	Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. <i>Carcinogenesis</i> , 2014, 35, 2723-2730.	2.8	57
18	A novel antisense long noncoding RNA regulates the expression of MDC1 in bladder cancer. <i>Oncotarget</i> , 2015, 6, 484-493.	1.8	56

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19	Genetic variants in noncoding PIWI-interacting RNA and colorectal cancer risk. <i>Cancer</i> , 2015, 121, 2044-2052.	4.1	56
20	Large-scale association analysis in Asians identifies new susceptibility loci for prostate cancer. <i>Nature Communications</i> , 2015, 6, 8469.	12.8	51
21	miR-107 regulates tumor progression by targeting NF1 in gastric cancer. <i>Scientific Reports</i> , 2016, 6, 36531.	3.3	51
22	Thymidylate Synthase 5' and 3'-Untranslated Region Polymorphisms Associated with Risk and Progression of Squamous Cell Carcinoma of the Head and Neck. <i>Clinical Cancer Research</i> , 2004, 10, 7903-7910.	7.0	47
23	Genetic variations in microRNAs and the risk and survival of renal cell cancer. <i>Carcinogenesis</i> , 2014, 35, 1629-1635.	2.8	47
24	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.	0.9	46
25	High-Level Genetic Diversity and Complex Population Structure of Siberian Apricot (<i>Prunus sibirica</i> L.) in China as Revealed by Nuclear SSR Markers. <i>PLoS ONE</i> , 2014, 9, e87381.	2.5	46
26	A functional variant in miR-143 promoter contributes to prostate cancer risk. <i>Archives of Toxicology</i> , 2016, 90, 403-414.	4.2	43
27	Ambient fine particulate matter (PM2.5) induces oxidative stress and pro-inflammatory response via up-regulating the expression of CYP1A1/1B1 in human bronchial epithelial cells in vitro. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2019, 839, 40-48.	1.7	42
28	An inverse association between tea consumption and colorectal cancer risk. <i>Oncotarget</i> , 2017, 8, 37367-37376.	1.8	42
29	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.	19.2	41
30	Short-term effects of ambient air pollution and childhood lower respiratory diseases. <i>Scientific Reports</i> , 2017, 7, 4414.	3.3	41
31	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.	7.0	39
32	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020, 41, 8-17.	2.8	38
33	A Functional Ser326Cys Polymorphism in hOGG1 Is Associated with Noise-Induced Hearing Loss in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e89662.	2.5	37
34	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.	5.3	37
35	The HOTAIR, PRNCR1 and POLR2E polymorphisms are associated with cancer risk: a meta-analysis. <i>Oncotarget</i> , 2017, 8, 43271-43283.	1.8	37
36	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.	12.1	36

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37	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.	1.0	35
38	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.	2.8	35
39	Body mass index (BMI) trajectories and risk of colorectal cancer in the PLCO cohort. <i>British Journal of Cancer</i> , 2018, 119, 130-132.	6.4	35
40	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.	0.8	34
41	Circulating MicroRNA-26a in Plasma and Its Potential Diagnostic Value in Gastric Cancer. <i>PLoS ONE</i> , 2016, 11, e0151345.	2.5	34
42	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.	2.5	33
43	HOTAIR rs7958904 polymorphism is associated with increased cervical cancer risk in a Chinese population. <i>Scientific Reports</i> , 2017, 7, 3144.	3.3	32
44	METTL3 regulates PM2.5-induced cell injury by targeting OSGIN1 in human airway epithelial cells. <i>Journal of Hazardous Materials</i> , 2021, 415, 125573.	12.4	32
45	Cumulative effect of genome-wide association study-identified genetic variants for bladder cancer. <i>International Journal of Cancer</i> , 2014, 135, 2653-2660.	5.1	31
46	Pri-miR-34b/c rs4938723 polymorphism contributes to acute lymphoblastic leukemia susceptibility in Chinese children. <i>Leukemia and Lymphoma</i> , 2016, 57, 1436-1441.	1.3	31
47	The biogenesis and biological function of PIWI-interacting RNA in cancer. <i>Journal of Hematology and Oncology</i> , 2021, 14, 93.	17.0	31
48	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018, 92, 2681-2689.	4.2	30
49	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.	1.0	28
50	Association between obesity and bladder cancer recurrence: A meta-analysis. <i>Clinica Chimica Acta</i> , 2018, 480, 41-46.	1.1	28
51	LncRNA <i>PCAT1</i> and its genetic variant rs1902432 are associated with prostate cancer risk. <i>Journal of Cancer</i> , 2018, 9, 1414-1420.	2.5	28
52	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.	5.1	27
53	Alternative splicing related genetic variants contribute to bladder cancer risk. <i>Molecular Carcinogenesis</i> , 2020, 59, 923-929.	2.7	27
54	Assessing the Effectiveness of Problem-Based Learning of Preventive Medicine Education in China. <i>Scientific Reports</i> , 2014, 4, 5126.	3.3	25

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55	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.	5.1	25
56	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.	4.1	25
57	Polymorphism rs2682818 in miR-618 is associated with colorectal cancer susceptibility in a Han Chinese population. <i>Cancer Medicine</i> , 2018, 7, 1194-1200.	2.8	24
58	Association study between XPG Asp1104His polymorphism and colorectal cancer risk in a Chinese population. <i>Scientific Reports</i> , 2014, 4, 6700.	3.3	23
59	The association of rs710886 in lncRNA PCAT1 with bladder cancer risk in a Chinese population. <i>Gene</i> , 2017, 627, 226-232.	2.2	23
60	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.	10.3	23
61	Effect of PM2.5 exposure on circulating fibrinogen and IL-6 levels: A systematic review and meta-analysis. <i>Chemosphere</i> , 2021, 271, 129565.	8.2	23
62	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.	5.5	23
63	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.	3.3	22
64	Rare variants in BRCA2 and CHEK2 are associated with the risk of urinary tract cancers. <i>Scientific Reports</i> , 2016, 6, 33542.	3.3	22
65	Mxd1 mediates hypoxia-induced cisplatin resistance in osteosarcoma cells by repression of the PTEN tumor suppressor gene. <i>Molecular Carcinogenesis</i> , 2017, 56, 2234-2244.	2.7	22
66	A functional polymorphism in <i>TFF1</i> promoter is associated with the risk and prognosis of gastric cancer. <i>International Journal of Cancer</i> , 2018, 142, 1805-1816.	5.1	22
67	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.	14.5	22
68	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.	11.2	21
69	The effects of particulate matters on allergic rhinitis in Nanjing, China. <i>Environmental Science and Pollution Research</i> , 2019, 26, 11452-11457.	5.3	20
70	Relationship between particulate matter exposure and female breast cancer incidence and mortality: a systematic review and meta-analysis. <i>International Archives of Occupational and Environmental Health</i> , 2021, 94, 191-201.	2.3	19
71	Integrative omics provide biological and clinical insights into acute respiratory distress syndrome. <i>Intensive Care Medicine</i> , 2021, 47, 761-771.	8.2	19
72	Fine particulate matter induces METTL3-mediated m6A modification of BIRC5 mRNA in bladder cancer. <i>Journal of Hazardous Materials</i> , 2022, 437, 129310.	12.4	19

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73	Variants in angiogenesis-related genes and the risk of clear cell renal cell carcinoma. <i>Mutagenesis</i> , 2014, 29, 419-425.	2.6	18
74	Functional POR A503V is associated with the risk of bladder cancer in a Chinese population. <i>Scientific Reports</i> , 2015, 5, 11751.	3.3	18
75	The rs767649 polymorphism in the promoter of miR-155 contributes to the decreased risk for cervical cancer in a Chinese population. <i>Gene</i> , 2016, 595, 109-114.	2.2	18
76	Genetic variants in N6-methyladenosine are associated with bladder cancer risk in the Chinese population. <i>Archives of Toxicology</i> , 2021, 95, 299-309.	4.2	18
77	Genetic variants in m6A regulators are associated with gastric cancer risk. <i>Archives of Toxicology</i> , 2021, 95, 1081-1088.	4.2	18
78	Genetic Polymorphisms in IGF-I and IGFBP-3 Are Associated with Prostate Cancer in the Chinese Population. <i>PLoS ONE</i> , 2014, 9, e85609.	2.5	18
79	SNORA74B gene silencing inhibits gallbladder cancer cells by inducing PHLPP and suppressing Akt/mTOR signaling. <i>Oncotarget</i> , 2017, 8, 19980-19996.	1.8	18
80	A genetic study and meta-analysis of the genetic predisposition of prostate cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 21393-21403.	1.8	18
81	Polymorphisms of the DNA repair gene MGMT and risk and progression of head and neck cancer. <i>DNA Repair</i> , 2010, 9, 558-566.	2.8	17
82	A novel mechanism of rs763110 polymorphism contributing to cervical cancer risk by affecting the binding affinity of C/EBP β and OCT1 complex to chromatin. <i>International Journal of Cancer</i> , 2017, 140, 756-763.	5.1	17
83	Long non-coding RNA FLJ22763 is involved in the progression and prognosis of gastric cancer. <i>Gene</i> , 2019, 693, 84-91.	2.2	17
84	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.	4.1	16
85	A genetic variant of miR-148a binding site in the SCRNI 3'UTR is associated with susceptibility and prognosis of gastric cancer. <i>Scientific Reports</i> , 2014, 4, 7080.	3.3	16
86	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020, 137, 105543.	10.0	16
87	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.	5.3	16
88	JWA, a novel microtubule-associated protein, regulates homeostasis of intracellular amino acids in PC12 cells. <i>Science Bulletin</i> , 2003, 48, 1828-1834.	1.7	15
89	Polymorphisms and haplotypes of serine hydroxymethyltransferase and risk of squamous cell carcinoma of the head and neck: a case-control analysis. <i>Pharmacogenetics and Genomics</i> , 2005, 15, 557-564.	1.5	15
90	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.	2.2	15

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91	A Genetic Variant Located in miR-146b Promoter Region Is Associated with Prognosis of Gastric Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 822-828.	2.5	15
92	Hypermethylation of EIF4E promoter is associated with early onset of gastric cancer. <i>Carcinogenesis</i> , 2018, 39, 66-71.	2.8	15
93	Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. <i>Journal of Gastroenterology</i> , 2019, 54, 939-949.	5.1	15
94	Photografting of unable-to-be-irradiated surfaces. I. Batch vapor-phase process by one-step method. <i>Journal of Applied Polymer Science</i> , 2006, 101, 2269-2276.	2.6	14
95	MDM2 SNP309 polymorphism is associated with colorectal cancer risk. <i>Scientific Reports</i> , 2014, 4, 4851.	3.3	14
96	PSCA rs2294008 polymorphism contributes to the decreased risk for cervical cancer in a Chinese population. <i>Scientific Reports</i> , 2016, 6, 23465.	3.3	14
97	Genetic variants in PI3K/Akt/mTOR pathway genes contribute to gastric cancer risk. <i>Gene</i> , 2018, 670, 130-135.	2.2	14
98	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.	4.2	14
99	MUC1 is associated with TFF2 methylation in gastric cancer. <i>Clinical Epigenetics</i> , 2020, 12, 37.	4.1	14
100	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.	10.0	14
101	Vitamin B2 intake reduces the risk for colorectal cancer: a dose-response analysis. <i>European Journal of Nutrition</i> , 2019, 58, 1591-1602.	3.9	13
102	Simultaneous quantification of five phenols in settled house dust using ultra-high performance liquid chromatography-tandem mass spectrometry. <i>Analytical Methods</i> , 2013, 5, 5339.	2.7	12
103	Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. <i>Journal of Cancer</i> , 2017, 8, 1355-1361.	2.5	12
104	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.	5.3	12
105	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018, 673, 174-180.	2.2	12
106	Genetic Variants in RKIP Are Associated with Clear Cell Renal Cell Carcinoma Risk in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e109285.	2.5	12
107	XRCC1 mediated the development of cervical cancer through a novel Sp1/Krox-20 switch. <i>Oncotarget</i> , 2017, 8, 86217-86226.	1.8	12
108	Functional annotation of colorectal cancer susceptibility loci identifies <i>MLH1</i> associated with MSI patients. <i>Gut</i> , 2016, 65, 1227-1228.	12.1	11

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109	Genetic variations in Hippo pathway genes influence bladder cancer risk in a Chinese population. Archives of Toxicology, 2020, 94, 785-794.	4.2	11
110	Photografting of unable-to-be-irradiated surfaces. II. Batch liquid-phase process by one-step method. Journal of Applied Polymer Science, 2007, 103, 118-124.	2.6	10
111	A miR-29c binding site genetic variant in the 3' untranslated region of LAMTOR3 gene is associated with gastric cancer risk. Biomedicine and Pharmacotherapy, 2015, 69, 70-75.	5.6	10
112	Identification of a novel susceptibility locus at 16q23.1 associated with childhood acute lymphoblastic leukemia in Han Chinese. Human Molecular Genetics, 2016, 25, ddw112.	2.9	10
113	Genetic variants, PM2.5 exposure level and global DNA methylation level: A multi-center population-based study in Chinese. Toxicology Letters, 2017, 269, 77-82.	0.8	10
114	Tagging SNPs in the HOTAIR gene are associated with bladder cancer risk in a Chinese population. Gene, 2018, 664, 22-26.	2.2	10
115	Genetic Variant in Long Non-Coding RNA H19 Modulates Its Expression and Predicts Renal Cell Carcinoma Susceptibility and Mortality. Frontiers in Oncology, 2020, 10, 785.	2.8	10
116	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.	5.1	9
117	Genetic variant in miR-21 binding sites is associated with colorectal cancer risk. Journal of Cellular and Molecular Medicine, 2019, 23, 2012-2019.	3.6	9
118	A MAP3k1 SNP Predicts Survival of Gastric Cancer in a Chinese Population. PLoS ONE, 2014, 9, e96083.	2.5	9
119	No Association between hOGG1 Ser326Cys polymorphism and risk of squamous cell carcinoma of the head and neck. Cancer Epidemiology Biomarkers and Prevention, 2004, 13, 1081-3.	2.5	9
120	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.	2.3	9
121	Associations of NR5A2 Gene Polymorphisms with the Clinicopathological Characteristics and Survival of Gastric Cancer. International Journal of Molecular Sciences, 2014, 15, 22902-22917.	4.1	8
122	Genetic variation in C12orf51 is associated with prognosis of intestinal-type gastric cancer in a Chinese population. Biomedicine and Pharmacotherapy, 2015, 69, 133-138.	5.6	8
123	The association analysis of hOGG1 genetic variants and gastric cancer risk in a Chinese population. Oncotarget, 2016, 7, 66061-66068.	1.8	8
124	Identification of low-frequency variants of UGT1A3 associated with bladder cancer risk by next-generation sequencing. Oncogene, 2021, 40, 2382-2394.	5.9	8
125	Global internet search trends related to gastrointestinal symptoms predict regional COVID-19 outbreaks. Journal of Infection, 2022, 84, 56-63.	3.3	8
126	Evaluation of genome-wide genotyping concordance between tumor tissues and peripheral blood. Genomics, 2017, 109, 108-112.	2.9	7

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127	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018, 33, 82-87.	6.1	7
128	Genetic variants in SMARC genes are associated with DNA damage levels in Chinese population. <i>Toxicology Letters</i> , 2014, 229, 327-332.	0.8	6
129	Genetic variation in IGF1 predicts renal cell carcinoma susceptibility and prognosis in Chinese population. <i>Scientific Reports</i> , 2016, 6, 39014.	3.3	6
130	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.	1.0	6
131	Information transduction capacity reduces the uncertainties in annotation-free isoform discovery and quantification. <i>Nucleic Acids Research</i> , 2017, 45, e143-e143.	14.5	6
132	Probabilistic natural mapping of gene-level tests for genome-wide association studies. <i>Briefings in Bioinformatics</i> , 2018, 19, 545-553.	6.5	6
133	Effects of TSP-1 -696 C/T polymorphism on bladder cancer susceptibility and clinicopathologic features. <i>Cancer Genetics</i> , 2014, 207, 247-252.	0.4	5
134	Novel CpG-SNPs in the gastric acid secretion pathway GNAI3 and susceptibility to gastric cancer. <i>Gene</i> , 2020, 736, 144447.	2.2	5
135	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.	4.2	5
136	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.	4.1	5
137	Genetic Variations in the 3'UTR-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.	2.0	5
138	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.	4.2	4
139	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021, 95, 2523-2532.	4.2	4
140	Association Between MIF-AS rs755622 and Nephrolithiasis Risk in a Chinese Population. <i>Medical Science Monitor</i> , 2016, 22, 563-568.	1.1	4
141	Mxi1-0 regulates the growth of human umbilical vein endothelial cells through extracellular signal-regulated kinase 1/2 (ERK1/2) and interleukin-8 (IL-8)-dependent pathways. <i>PLoS ONE</i> , 2017, 12, e0178831.	2.5	4
142	Functional variants of RPS6KB1 and PIK3R1 in the autophagy pathway genes and risk of bladder cancer. <i>Archives of Toxicology</i> , 2021, , 1.	4.2	4
143	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, , .	2.9	4
144	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.	1.6	3

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145	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.	1.0	3
146	Genetic variants in XDH are associated with prognosis for gastric cancer in a Chinese population. Gene, 2018, 663, 196-202.	2.2	3
147	Polymorphism rs4787951 in IL-4R contributes to the increased risk of renal cell carcinoma in a Chinese population. Gene, 2019, 685, 242-247.	2.2	3
148	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. Archives of Toxicology, 2020, 94, 325-333.	4.2	3
149	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.	3.6	3
150	Identification of common genetic variants associated with serum concentrations of p, p'-DDE in non-occupational populations in eastern China. Environment International, 2021, 152, 106507.	10.0	3
151	RPTOR methylation in the peripheral blood and breast cancer in the Chinese population. Genes and Genomics, 2021, , 1.	1.4	3
152	Genome-Wide Association Analyses Identify <i>CATSPERE</i> as a Mediator of Colorectal Cancer Susceptibility and Progression. Cancer Research, 2022, 82, 986-997.	0.9	3
153	Polymorphism of methylenetetrahydrofolate reductase gene is associated with response to fluorouracil-based chemotherapy in Chinese patients with gastric cancer. Chinese Medical Journal, 2014, 127, 3562-7.	2.3	3
154	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.	4.2	2
155	Genetic variants in splicing factor genes and susceptibility to bladder cancer. Gene, 2022, 809, 146022.	2.2	1
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157	Comprehensive genetic mutation analysis of human gastric adenocarcinomas.. Journal of Clinical Oncology, 2013, 31, 4106-4106.	1.6	0
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