

Hongbing Shen

List of Publications by Year in descending order

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Version: 2024-02-01

439
papers

22,684
citations

12597

71
h-index

17891

125
g-index

454
all docs

454
docs citations

454
times ranked

33432
citing authors

#	ARTICLE	IF	CITATIONS
1	Risk Assessment for Longitudinal Trajectories of Modifiable Lifestyle Factors on Chronic Kidney Disease Burden in China: A Population-based Study. <i>Journal of Epidemiology</i> , 2022, 32, 449-455.	1.1	2
2	Identification of A-to-I RNA editing profiles and their clinical relevance in lung adenocarcinoma. <i>Science China Life Sciences</i> , 2022, 65, 19-32.	2.3	6
3	Comprehensive estimation for the length and dispersion of COVID-19 incubation period: a systematic review and meta-analysis. <i>Infection</i> , 2022, 50, 803-813.	2.3	12
4	Circulating C-reactive protein increases lung cancer risk: Results from a prospective cohort of UK Biobank. <i>International Journal of Cancer</i> , 2022, 150, 47-55.	2.3	15
5	Global COVID-19 Pandemic Waves: Limited Lessons Learned Worldwide over the Past Year. <i>Engineering</i> , 2022, 13, 91-98.	3.2	10
6	Adherence to Healthy Lifestyle and Liver cancer in Chinese: a prospective cohort study of 0.5 million people. <i>British Journal of Cancer</i> , 2022, 126, 815-821.	2.9	15
7	Accounting for EGFR Mutations in Epidemiologic Analyses of Non-Small Cell Lung Cancers: Examples Based on the International Lung Cancer Consortium Data. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 679-687.	1.1	1
8	Plasma metabolomic profiles for colorectal cancer precursors in women. <i>European Journal of Epidemiology</i> , 2022, 37, 413-422.	2.5	11
9	Genome-wide interaction analysis identified low-frequency variants with sex disparity in lung cancer risk. <i>Human Molecular Genetics</i> , 2022, 31, 2831-2843.	1.4	4
10	Genome-Wide Association Analyses Identify CATSPER1 as a Mediator of Colorectal Cancer Susceptibility and Progression. <i>Cancer Research</i> , 2022, 82, 986-997.	0.4	3
11	FSTL1 promotes liver fibrosis by reprogramming macrophage function through modulating the intracellular function of PKM2. <i>Gut</i> , 2022, 71, 2539-2550.	6.1	55
12	A polygenic risk score improves risk stratification of coronary artery disease: a large-scale prospective Chinese cohort study. <i>European Heart Journal</i> , 2022, 43, 1702-1711.	1.0	58
13	One-off low-dose CT for lung cancer screening in China: a multicentre, population-based, prospective cohort study. <i>Lancet Respiratory Medicine</i> , 2022, 10, 378-391.	5.2	69
14	Associations of systemic inflammation markers with identification of pulmonary nodule and incident lung cancer in Chinese population. <i>Cancer Medicine</i> , 2022, 11, 2482-2491.	1.3	13
15	Study on the Influencing Factors of Knowledge, Attitudes and Practice About Tuberculosis Among Freshmen in Jiangsu, China: A Cross-Sectional Study. <i>Infection and Drug Resistance</i> , 2022, Volume 15, 1235-1245.	1.1	4
16	A polygenic risk score for nasopharyngeal carcinoma shows potential for risk stratification and personalized screening. <i>Nature Communications</i> , 2022, 13, 1966.	5.8	19
17	Reply to: Contribution of COPD as a Mediator for the Association Between Air Pollution and Lung Cancer. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2022, , .	2.5	0
18	A Large-Scale Genome-Wide Gene-Gene Interaction Study of Lung Cancer Susceptibility in Europeans With a Trans-Ethnic Validation in Asians. <i>Journal of Thoracic Oncology</i> , 2022, 17, 974-990.	0.5	18

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19	Transcriptome-wide association study identifies <i>PSMB9</i> as a susceptibility gene for coal workers' pneumoconiosis. <i>Environmental Toxicology</i> , 2022, , .	2.1	1
20	Rapid Visual Detection of Pathogenic <i>Streptococcus suis</i> Type 2 through a Recombinase Polymerase Amplification Assay Coupled with Lateral Flow Test. <i>Zoonoses</i> , 2022, 2, .	0.5	0
21	Functional characterization and clinical significance of super-enhancers in lung adenocarcinoma. <i>Molecular Carcinogenesis</i> , 2022, 61, 776-786.	1.3	2
22	The genomic landscape of cholangiocarcinoma reveals the disruption of post-transcriptional modifiers. <i>Nature Communications</i> , 2022, 13, .	5.8	17
23	Associations of genetic risk, BMI trajectories, and the risk of non-small cell lung cancer: a population-based cohort study. <i>BMC Medicine</i> , 2022, 20, .	2.3	10
24	The continuing evolution of birth cohort studies: achievements and challenges. <i>Biology of Reproduction</i> , 2022, 107, 358-367.	1.2	1
25	A Lipid Signature with Perturbed Triacylglycerol Co-Regulation, Identified from Targeted Lipidomics, Predicts Risk for Type 2 Diabetes and Mediates the Risk from Adiposity in Two Prospective Cohorts of Chinese Adults. <i>Clinical Chemistry</i> , 2022, 68, 1094-1107.	1.5	3
26	Human X chromosome exome sequencing identifies <i>BCORL1</i> as contributor to spermatogenesis. <i>Journal of Medical Genetics</i> , 2021, 58, 56-65.	1.5	13
27	Integration of multiomic annotation data to prioritize and characterize inflammation and immune-related risk variants in squamous cell lung cancer. <i>Genetic Epidemiology</i> , 2021, 45, 99-114.	0.6	7
28	Hormonal factors in association with lung cancer among Asian women: A pooled analysis from the International Lung Cancer Consortium. <i>International Journal of Cancer</i> , 2021, 148, 2241-2254.	2.3	9
29	Causal relationships between body mass index, smoking and lung cancer: Univariable and multivariable Mendelian randomization. <i>International Journal of Cancer</i> , 2021, 148, 1077-1086.	2.3	73
30	Comprehensive functional annotation of susceptibility variants identifies genetic heterogeneity between lung adenocarcinoma and squamous cell carcinoma. <i>Frontiers of Medicine</i> , 2021, 15, 275-291.	1.5	21
31	Association between dietary sodium, potassium intake and lung cancer risk: evidence from the prostate, lung, colorectal and ovarian cancer screening trial and the Women's Health Initiative. <i>Translational Lung Cancer Research</i> , 2021, 10, 45-56.	1.3	16
32	Pre-diagnostic circulating concentrations of insulin-like growth factor-1 and risk of COVID-19 mortality: results from UK Biobank. <i>European Journal of Epidemiology</i> , 2021, 36, 311-318.	2.5	19
33	Sub-multiplicative interaction between polygenic risk score and household coal use in relation to lung adenocarcinoma among never-smoking women in Asia. <i>Environment International</i> , 2021, 147, 105975.	4.8	12
34	Assisted reproductive technology and birth defects in a Chinese birth cohort study. <i>The Lancet Regional Health - Western Pacific</i> , 2021, 7, 100090.	1.3	24
35	Prediction and clinical utility of a liver cancer risk model in Chinese adults: A prospective cohort study of 0.5 million people. <i>International Journal of Cancer</i> , 2021, 148, 2924-2934.	2.3	6
36	Antibody seroprevalence in the epicenter Wuhan, Hubei, and six selected provinces after containment of the first epidemic wave of COVID-19 in China. <i>The Lancet Regional Health - Western Pacific</i> , 2021, 8, 100094.	1.3	41

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37	Genetic variants associated with expression of TCF19 contribute to the risk of head and neck cancer in Chinese population. <i>Journal of Medical Genetics</i> , 2021, , jmedgenet-2020-107410.	1.5	7
38	Relationships between sleep traits and lung cancer risk: a prospective cohort study in UK Biobank. <i>Sleep</i> , 2021, 44, .	0.6	26
39	Child marriage, maternal serum metal exposure, and risk of preterm birth in rural Bangladesh: evidence from mediation analysis. <i>Journal of Exposure Science and Environmental Epidemiology</i> , 2021, 31, 571-580.	1.8	12
40	Comprehensive analyses of m6A regulators and interactive coding and non-coding RNAs across 32 cancer types. <i>Molecular Cancer</i> , 2021, 20, 67.	7.9	65
41	A cross-tissue transcriptome-wide association study identifies novel susceptibility genes for lung cancer in Chinese populations. <i>Human Molecular Genetics</i> , 2021, 30, 1666-1676.	1.4	9
42	Potential functional variants of KIAA genes are associated with breast cancer risk in a case control study. <i>Annals of Translational Medicine</i> , 2021, 9, 549-549.	0.7	1
43	Sex-specific associations of circulating testosterone levels with all-cause and cause-specific mortality. <i>European Journal of Endocrinology</i> , 2021, 184, 723-732.	1.9	17
44	A multi-omics study links TNS3 and SEPT7 to long-term former smoking NSCLC survival. <i>Npj Precision Oncology</i> , 2021, 5, 39.	2.3	9
45	Association of assisted reproductive technology, germline de novo mutations and congenital heart defects in a prospective birth cohort study. <i>Cell Research</i> , 2021, 31, 919-928.	5.7	26
46	An Improved Detection of Circulating Tumor DNA in Extracellular Vesicles-Depleted Plasma. <i>Frontiers in Oncology</i> , 2021, 11, 691798.	1.3	3
47	Sex Hormone and Colorectal Cancer: The Knowns and Unknowns. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 1302-1304.	1.1	7
48	Genome-wide gene-smoking interaction study identified novel susceptibility loci for non-small cell lung cancer in Chinese populations. <i>Carcinogenesis</i> , 2021, 42, 1154-1161.	1.3	1
49	Genetic Risk for Overall Cancer and the Benefit of Adherence to a Healthy Lifestyle. <i>Cancer Research</i> , 2021, 81, 4618-4627.	0.4	48
50	Comparative Characterization and Risk Stratification of Asymptomatic and Presymptomatic Patients With COVID-19. <i>Frontiers in Immunology</i> , 2021, 12, 700449.	2.2	2
51	Air Pollution, Genetic Factors, and the Risk of Lung Cancer: A Prospective Study in the UK Biobank. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021, 204, 817-825.	2.5	133
52	Association of maternal diabetes during pregnancy with high refractive error in offspring: a nationwide population-based cohort study. <i>Diabetologia</i> , 2021, 64, 2466-2477.	2.9	15
53	Diet and Risk of Incident Lung Cancer: A Large Prospective Cohort Study in UK Biobank. <i>American Journal of Clinical Nutrition</i> , 2021, 114, 2043-2051.	2.2	38
54	The rates and medical necessity of cesarean delivery in China, 2012-2019: an inspiration from Jiangsu. <i>BMC Medicine</i> , 2021, 19, 14.	2.3	8

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55	Plasma sex hormones and risk of conventional and serrated precursors of colorectal cancer in postmenopausal women. <i>BMC Medicine</i> , 2021, 19, 18.	2.3	12
56	The cancer-testis gene, <i>MEIOB</i> , sensitizes triple-negative breast cancer to PARP1 inhibitors by inducing homologous recombination deficiency. <i>Cancer Biology and Medicine</i> , 2021, 18, 74-87.	1.4	7
57	Stress, anxiety, and depression in infertile couples are not associated with a first IVF or ICSI treatment outcome. <i>BMC Pregnancy and Childbirth</i> , 2021, 21, 725.	0.9	6
58	Development of Rapid and Visual Nucleic Acid Detection Methods towards Four Serotypes of Human Adenovirus Species B Based on RPA-LF Test. <i>BioMed Research International</i> , 2021, 2021, 1-13.	0.9	3
59	Bi-allelic variants in human <i>WDR63</i> cause male infertility via abnormal inner dynein arms assembly. <i>Cell Discovery</i> , 2021, 7, 110.	3.1	19
60	Polygenic risk scores: the future of cancer risk prediction, screening, and precision prevention. <i>Medical Review</i> , 2021, 1, 129-149.	0.3	4
61	Tea consumption and risk of stroke in Chinese adults: a prospective cohort study of 0.5 million men and women. <i>American Journal of Clinical Nutrition</i> , 2020, 111, 197-206.	2.2	27
62	Cancer incidence in relation to body fatness among 0.5 million men and women: Findings from the China Kadoorie Biobank. <i>International Journal of Cancer</i> , 2020, 146, 987-998.	2.3	19
63	Colorectal cancer susceptibility variants and risk of conventional adenomas and serrated polyps: results from three cohort studies. <i>International Journal of Epidemiology</i> , 2020, 49, 259-269.	0.9	13
64	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
65	Tuberculosis infection and lung adenocarcinoma: Mendelian randomization and pathway analysis of genome-wide association study data from never-smoking Asian women. <i>Genomics</i> , 2020, 112, 1223-1232.	1.3	15
66	Genome-wide association study of INDELs identified four novel susceptibility loci associated with lung cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2855-2864.	2.3	7
67	Genetic risk of extranodal natural killer T-cell lymphoma: a genome-wide association study in multiple populations. <i>Lancet Oncology</i> , 2020, 21, 306-316.	5.1	49
68	Association Between Levels of Sex Hormones and Risk of Esophageal Adenocarcinoma and Barrett's Esophagus. <i>Clinical Gastroenterology and Hepatology</i> , 2020, 18, 2701-2709.e3.	2.4	12
69	Genetic risk, incident gastric cancer, and healthy lifestyle: a meta-analysis of genome-wide association studies and prospective cohort study. <i>Lancet Oncology</i> , 2020, 21, 1378-1386.	5.1	123
70	Comprehensive characterization of functional eRNAs in lung adenocarcinoma reveals novel regulators and a prognosis-related molecular subtype. <i>Theranostics</i> , 2020, 10, 11264-11277.	4.6	20
71	Analysis of the interaction effect of 48 SNPs and obesity on type 2 diabetes in Chinese Hans. <i>BMJ Open Diabetes Research and Care</i> , 2020, 8, e001638.	1.2	4
72	Tumor evolutionary trajectories during the acquisition of invasiveness in early stage lung adenocarcinoma. <i>Nature Communications</i> , 2020, 11, 6083.	5.8	15

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73	Alterations of gut microbiota composition in neonates conceived by assisted reproductive technology and its relation to infant growth. <i>Gut Microbes</i> , 2020, 12, 1794466.	4.3	9
74	Metabolomic Signatures of Long-term Coffee Consumption and Risk of Type 2 Diabetes in Women. <i>Diabetes Care</i> , 2020, 43, 2588-2596.	4.3	27
75	Identification of Recurrent Variants in BRCA1 and BRCA2 across Multiple Cancers in the Chinese Population. <i>BioMed Research International</i> , 2020, 2020, 1-11.	0.9	4
76	Effect of endometrial thickness and embryo quality on live-birth rate of fresh IVF/ICSI cycles: a retrospective cohort study. <i>Reproductive Biology and Endocrinology</i> , 2020, 18, 89.	1.4	18
77	Implementation of Clinical Diagnostic Criteria and Universal Symptom Survey Contributed to Lower Magnitude and Faster Resolution of the COVID-19 Epidemic in Wuhan. <i>Engineering</i> , 2020, 6, 1141-1146.	3.2	14
78	Physical activity and health in Chinese children and adolescents: expert consensus statement (2020). <i>British Journal of Sports Medicine</i> , 2020, 54, 1321-1331.	3.1	71
79	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
80	Transcriptome-wide association study for persistent hepatitis B virus infection and related hepatocellular carcinoma. <i>Liver International</i> , 2020, 40, 2117-2127.	1.9	6
81	Spatial statistical analysis of Coronavirus Disease 2019 (Covid-19) in China. <i>Geospatial Health</i> , 2020, 15, .	0.3	36
82	Clinical characteristics of 24 asymptomatic infections with COVID-19 screened among close contacts in Nanjing, China. <i>Science China Life Sciences</i> , 2020, 63, 706-711.	2.3	1,090
83	Systematic analysis of genetic variants in cancer-testis genes identified two novel lung cancer susceptibility loci in Chinese population. <i>Journal of Cancer</i> , 2020, 11, 1985-1993.	1.2	2
84	Possible Transmission of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) in a Public Bath Center in Huai'an, Jiangsu Province, China. <i>JAMA Network Open</i> , 2020, 3, e204583.	2.8	85
85	Vitamin D Status and Risk of All-Cause and Cause-Specific Mortality in a Large Cohort: Results From the UK Biobank. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, e3606-e3619.	1.8	60
86	Systematic analyses of genetic variants in chromatin interaction regions identified four novel lung cancer susceptibility loci. <i>Journal of Cancer</i> , 2020, 11, 1075-1081.	1.2	16
87	Genetic Variations in miR-30 Family Member Regulatory Regions Are Associated with Breast Cancer Risk in a Chinese Population. <i>BioMed Research International</i> , 2020, 2020, 1-8.	0.9	1
88	Association Analysis of Driver Gene-Related Genetic Variants Identified Novel Lung Cancer Susceptibility Loci with 20,871 Lung Cancer Cases and 15,971 Controls. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1423-1429.	1.1	6
89	From human genome epidemiology to systems epidemiology: current progress and future perspective. <i>Journal of Biomedical Research</i> , 2020, 34, 323.	0.7	0
90	Tobacco smoking, alcohol drinking, betel quid chewing, and the risk of head and neck cancer in an East Asian population. <i>Head and Neck</i> , 2019, 41, 92-102.	0.9	63

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91	Identification of risk loci and a polygenic risk score for lung cancer: a large-scale prospective cohort study in Chinese populations. <i>Lancet Respiratory Medicine</i> , 2019, 7, 881-891.	5.2	167
92	Genomic signatures reveal DNA damage response deficiency in colorectal cancer brain metastases. <i>Nature Communications</i> , 2019, 10, 3190.	5.8	64
93	Fine Mapping in Chromosome 3q28 Identified Two Variants Associated with Lung Cancer Risk in Asian Population. <i>Journal of Cancer</i> , 2019, 10, 1862-1869.	1.2	1
94	Plasma Biomarkers of Insulin and the Insulin-like Growth Factor Axis, and Risk of Colorectal Adenoma and Serrated Polyp. <i>JNCI Cancer Spectrum</i> , 2019, 3, pkz056.	1.4	9
95	Mendelian randomization study of telomere length and lung cancer risk in East Asian population. <i>Cancer Medicine</i> , 2019, 8, 7469-7476.	1.3	15
96	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
97	Genome-wide analysis of expression quantitative trait loci identified potential lung cancer susceptibility variants among Asian populations. <i>Carcinogenesis</i> , 2019, 40, 263-268.	1.3	3
98	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 935-942.	1.1	21
99	The inherited variations of a p53-responsive enhancer in 13q12.12 confer lung cancer risk by attenuating TNFRSF19 expression. <i>Genome Biology</i> , 2019, 20, 103.	3.8	27
100	Identification of Novel T1D Risk Loci and Their Association With Age and Islet Function at Diagnosis in Autoantibody-Positive T1D Individuals: Based on a Two-Stage Genome-Wide Association Study. <i>Diabetes Care</i> , 2019, 42, 1414-1421.	4.3	60
101	Leukocyte telomere length, lipid parameters and gestational diabetes risk: a case-control study in a Chinese population. <i>Scientific Reports</i> , 2019, 9, 8483.	1.6	10
102	Associations Between Hepatitis B Virus Infection and Risk of All Cancer Types. <i>JAMA Network Open</i> , 2019, 2, e195718.	2.8	114
103	Body mass index and the risk of head and neck cancer in the Chinese population. <i>Cancer Epidemiology</i> , 2019, 60, 208-215.	0.8	14
104	Precision oncology of lung cancer: genetic and genomic differences in Chinese population. <i>Npj Precision Oncology</i> , 2019, 3, 14.	2.3	31
105	Association of expression quantitative trait loci for long noncoding RNAs with lung cancer risk in Asians. <i>Molecular Carcinogenesis</i> , 2019, 58, 1303-1313.	1.3	6
106	Hypomethylation-mediated activation of cancer/testis antigen KIF20A facilitates hepatocellular carcinoma progression through activating the Notch1/Hes1 signalling. <i>Cell Proliferation</i> , 2019, 52, e12581.	2.4	25
107	Coffee consumption and plasma biomarkers of metabolic and inflammatory pathways in US health professionals. <i>American Journal of Clinical Nutrition</i> , 2019, 109, 635-647.	2.2	59
108	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

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109	Commentary: Premalignant genomic data tracing the evolution of lung adenocarcinoma. <i>EBioMedicine</i> , 2019, 42, 16-17.	2.7	0
110	Blood groups A and AB are associated with increased gastric cancer risk: evidence from a large genetic study and systematic review. <i>BMC Cancer</i> , 2019, 19, 164.	1.1	30
111	Involuntary smoking and the risk of head and neck cancer in an East Asian population. <i>Cancer Epidemiology</i> , 2019, 59, 173-177.	0.8	8
112	A nomogram to predict overall survival of patients with early stage non-small cell lung cancer. <i>Journal of Thoracic Disease</i> , 2019, 11, 5407-5416.	0.6	16
113	5-Hydroxytryptamine Receptor 1D Aggravates Hepatocellular Carcinoma Progression Through FoxO6 in AKT-Dependent and Independent Manners. <i>Hepatology</i> , 2019, 69, 2031-2047.	3.6	33
114	Association of Mosaic Loss of Chromosome Y with Lung Cancer Risk and Prognosis in a Chinese Population. <i>Journal of Thoracic Oncology</i> , 2019, 14, 37-44.	0.5	19
115	Systematic analyses of regulatory variants in DNase I hypersensitive sites identified two novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2019, 40, 432-440.	1.3	5
116	A cancer-testis non-coding RNA LIN28B-AS1 activates driver gene LIN28B by interacting with IGF2BP1 in lung adenocarcinoma. <i>Oncogene</i> , 2019, 38, 1611-1624.	2.6	61
117	Spontaneous Seroclearance of Hepatitis B Surface Antigen and Risk of Hepatocellular Carcinoma. <i>Clinical Gastroenterology and Hepatology</i> , 2019, 17, 1204-1206.	2.4	14
118	Cross-Cancer Pleiotropic Analysis Reveals Novel Susceptibility Loci for Lung Cancer. <i>Frontiers in Oncology</i> , 2019, 9, 1492.	1.3	6
119	Interaction analysis between germline susceptibility loci and somatic alterations in lung cancer. <i>International Journal of Cancer</i> , 2018, 143, 878-885.	2.3	13
120	Association of 48 type 2 diabetes susceptibility loci with fasting plasma glucose and lipid levels in Chinese Hans. <i>Diabetes Research and Clinical Practice</i> , 2018, 139, 114-121.	1.1	4
121	Longitudinal associations of lifetime adiposity with leukocyte telomere length and mitochondrial DNA copy number. <i>European Journal of Epidemiology</i> , 2018, 33, 485-495.	2.5	28
122	miR-142-3p regulates autophagy by targeting ATG16L1 in thymic-derived regulatory T cell (tTreg). <i>Cell Death and Disease</i> , 2018, 9, 290.	2.7	37
123	Evaluation of CpG-SNPs in miRNA promoters and risk of breast cancer. <i>Gene</i> , 2018, 651, 1-8.	1.0	15
124	Association Analysis Identifies New Risk Loci for Coal Workers' Pneumoconiosis in Han Chinese Men. <i>Toxicological Sciences</i> , 2018, 163, 206-213.	1.4	14
125	Genome-Wide Association Study Identifies a New Locus at 7q21.13 Associated with Hepatitis B Virus-Related Hepatocellular Carcinoma. <i>Clinical Cancer Research</i> , 2018, 24, 906-915.	3.2	37
126	A novel plasma circular RNA circFARSA is a potential biomarker for non-small cell lung cancer. <i>Cancer Medicine</i> , 2018, 7, 2783-2791.	1.3	167

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127	PRDM8 exhibits antitumor activities toward hepatocellular carcinoma by targeting NAP1L1. <i>Hepatology</i> , 2018, 68, 994-1009.	3.6	35
128	Predicting the outbreak of hand, foot, and mouth disease in Nanjing, China: a time-series model based on weather variability. <i>International Journal of Biometeorology</i> , 2018, 62, 565-574.	1.3	24
129	Integrating expression-related SNPs into genome-wide gene and pathway-based analyses identified novel lung cancer susceptibility genes. <i>International Journal of Cancer</i> , 2018, 142, 1602-1610.	2.3	14
130	Low-dose CT for lung cancer screening: opportunities and challenges. <i>Frontiers of Medicine</i> , 2018, 12, 116-121.	1.5	28
131	Association of <i>CETP</i> Gene Variants With Risk for Vascular and Nonvascular Diseases Among Chinese Adults. <i>JAMA Cardiology</i> , 2018, 3, 34.	3.0	54
132	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. <i>Nature Communications</i> , 2018, 9, 3927.	5.8	43
133	Fine mapping in <i>TERT</i> region identified three independent lung cancer susceptibility signals: A large-scale multi-ethnic population study. <i>Molecular Carcinogenesis</i> , 2018, 57, 1289-1299.	1.3	5
134	Whole-genome sequencing reveals genomic signatures associated with the inflammatory microenvironments in Chinese NSCLC patients. <i>Nature Communications</i> , 2018, 9, 2054.	5.8	68
135	Genetically determined height was associated with lung cancer risk in East Asian population. <i>Cancer Medicine</i> , 2018, 7, 3445-3452.	1.3	6
136	Dataset for regulation between lncRNAs and their nearby protein-coding genes in human cancers. <i>Data in Brief</i> , 2018, 19, 1902-1906.	0.5	11
137	Spermine Alleviates Acute Liver Injury by Inhibiting Liver-Resident Macrophage Pro-Inflammatory Response Through ATG5-Dependent Autophagy. <i>Frontiers in Immunology</i> , 2018, 9, 948.	2.2	65
138	Tea, coffee, and head and neck cancer risk in a multicenter study in east Asia. <i>Oral Cancer</i> , 2018, 2, 57-65.	0.3	1
139	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018, 9, 3221.	5.8	60
140	Influencing factors of pregnancy loss and survival probability of clinical pregnancies conceived through assisted reproductive technology. <i>Reproductive Biology and Endocrinology</i> , 2018, 16, 74.	1.4	33
141	Systematic analysis reveals long noncoding RNAs regulating neighboring transcription factors in human cancers. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018, 1864, 2785-2792.	1.8	22
142	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
143	Exome-Wide Association Study Identified New Risk Loci for Hirschsprung's Disease. <i>Molecular Neurobiology</i> , 2017, 54, 1777-1785.	1.9	12
144	<i>KIT</i> polymorphisms were associated with the risk for head and neck squamous carcinoma in Chinese population. <i>Molecular Carcinogenesis</i> , 2017, 56, 232-237.	1.3	4

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145	Effects of potentially functional polymorphisms in suppressor of cytokine signaling 3 (<i>SOCS3</i>) on the risk of head and neck squamous cancer. <i>Journal of Oral Pathology and Medicine</i> , 2017, 46, 598-602.	1.4	6
146	Exome-Wide Association Study Identifies Low-Frequency Coding Variants in 2p23.2 and 7p11.2 Associated with Survival of Non-Small Cell Lung Cancer Patients. <i>Journal of Thoracic Oncology</i> , 2017, 12, 644-656.	0.5	10
147	The LINK-A lncRNA interacts with PtdIns(3,4,5)P3 to hyperactivate AKT and confer resistance to AKT inhibitors. <i>Nature Cell Biology</i> , 2017, 19, 238-251.	4.6	201
148	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
149	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
150	Potentially functional variants in lncRNAs are associated with breast cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2017, 56, 2048-2057.	1.3	11
151	Comprehensive assessment showed no associations of variants at the SLC10A1 locus with susceptibility to persistent HBV infection among Southern Chinese. <i>Scientific Reports</i> , 2017, 7, 46490.	1.6	20
152	A polymorphism in miR-1262 regulatory region confers the risk of lung cancer in Chinese population. <i>International Journal of Cancer</i> , 2017, 141, 958-966.	2.3	26
153	Genetic variants in autophagy associated genes are associated with DNA damage levels in Chinese population. <i>Gene</i> , 2017, 626, 414-419.	1.0	0
154	Genetically predicted high body mass index is associated with increased gastric cancer risk. <i>European Journal of Human Genetics</i> , 2017, 25, 1061-1066.	1.4	20
155	Cancer incidence and mortality: A cohort study in China, 2008-2013. <i>International Journal of Cancer</i> , 2017, 141, 1315-1323.	2.3	124
156	Race and Sex Differences of Long-Term Blood Pressure Profiles From Childhood and Adult Hypertension. <i>Hypertension</i> , 2017, 70, 66-74.	1.3	84
157	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017, 49, 1126-1132.	9.4	472
158	Targeted sequencing of chromosome 15q25 identified novel variants associated with risk of lung cancer and smoking behavior in Chinese. <i>Carcinogenesis</i> , 2017, 38, 552-558.	1.3	10
159	Genetic variants at 9p21.3 are associated with risk of esophageal squamous cell carcinoma in a Chinese population. <i>Cancer Science</i> , 2017, 108, 250-255.	1.7	14
160	Fine mapping of chromosome 5p15.33 identifies novel lung cancer susceptibility loci in Han Chinese. <i>International Journal of Cancer</i> , 2017, 141, 447-456.	2.3	17
161	Systematical analyses of variants in DNase I hypersensitive sites to identify hepatocellular carcinoma susceptibility loci in a Chinese population. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 1887-1894.	1.4	2
162	Diet and the risk of head-and-neck cancer among never-smokers and smokers in a Chinese population. <i>Cancer Epidemiology</i> , 2017, 46, 20-26.	0.8	15

#	ARTICLE	IF	CITATIONS
163	Impact of oral hygiene on head and neck cancer risk in a Chinese population. <i>Head and Neck</i> , 2017, 39, 2549-2557.	0.9	17
164	Metabolome-wide association study identified the association between a circulating polyunsaturated fatty acids variant rs174548 and lung cancer. <i>Carcinogenesis</i> , 2017, 38, 1147-1154.	1.3	21
165	Fine-mapping the MHC region in Asian populations identified novel variants modifying susceptibility to lung cancer. <i>Lung Cancer</i> , 2017, 112, 169-175.	0.9	21
166	The sex ratio of singleton and twin delivery offspring in assisted reproductive technology in China. <i>Scientific Reports</i> , 2017, 7, 7754.	1.6	18
167	A cis-eQTL genetic variant of the cancer-associated testis gene <i>CCDC116</i> is associated with risk of multiple cancers. <i>Human Genetics</i> , 2017, 136, 987-997.	1.8	7
168	Independent prognostic role of human papillomavirus genotype in cervical cancer. <i>BMC Infectious Diseases</i> , 2017, 17, 391.	1.3	43
169	Low-frequency nonsynonymous variants in <i>FKBP1</i> and <i>ARPC1B</i> genes are associated with breast cancer risk in Chinese women. <i>Molecular Carcinogenesis</i> , 2017, 56, 774-780.	1.3	12
170	Genetic variants affecting telomere length are associated with the prognosis of esophageal squamous cell carcinoma in a Chinese population. <i>Molecular Carcinogenesis</i> , 2017, 56, 1021-1029.	1.3	7
171	Estimation of heritability for nine common cancers using data from genome-wide association studies in Chinese population. <i>International Journal of Cancer</i> , 2017, 140, 329-336.	2.3	66
172	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 126-135.	1.1	278
173	Genetic Variants in the Promoter Region of <i>miR-10b</i> and the Risk of Breast Cancer. <i>BioMed Research International</i> , 2017, 2017, 1-7.	0.9	7
174	Human papillomavirus in semen and the risk for male infertility: a systematic review and meta-analysis. <i>BMC Infectious Diseases</i> , 2017, 17, 714.	1.3	80
175	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017, 12, e0177875.	1.1	79
176	Risk assessment models for genetic risk predictors of lung cancer using two-stage replication for Asian and European populations. <i>Oncotarget</i> , 2017, 8, 53959-53967.	0.8	11
177	Persistence of type-specific human papillomavirus infection among Daqing City women in China with normal cytology: a pilot prospective study. <i>Oncotarget</i> , 2017, 8, 81455-81461.	0.8	10
178	Systematic identification of long non-coding RNAs with cancer-testis expression patterns in 14 cancer types. <i>Oncotarget</i> , 2017, 8, 94769-94779.	0.8	10
179	Family History and Stroke Risk in China: Evidence from a Large Cohort Study. <i>Journal of Stroke</i> , 2017, 19, 188-195.	1.4	21
180	Association of <i>KCTD10</i> , <i>MVK</i> , and <i>MMAB</i> polymorphisms with dyslipidemia and coronary heart disease in Han Chinese population. <i>Lipids in Health and Disease</i> , 2016, 15, 171.	1.2	18

#	ARTICLE	IF	CITATIONS
181	Telomere length, genetic variants and risk of squamous cell carcinoma of the head and neck in Southeast Chinese. <i>Scientific Reports</i> , 2016, 6, 20675.	1.6	13
182	Evaluation of common variants in MG53 and the risk of type 2 diabetes and insulin resistance in Han Chinese. <i>SpringerPlus</i> , 2016, 5, 612.	1.2	5
183	The known genetic loci for telomere length may be involved in the modification of telomeres length after birth. <i>Scientific Reports</i> , 2016, 6, 38729.	1.6	17
184	Association between GWAS-identified lung adenocarcinoma susceptibility loci and EGFR mutations in never-smoking Asian women, and comparison with findings from Western populations. <i>Human Molecular Genetics</i> , 2016, 26, ddw414.	1.4	50
185	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016, 7, 11478.	5.8	73
186	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
187	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
188	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
189	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016, 76, 5103-5114.	0.4	100
190	Plasma metabolomics identified novel metabolites associated with risk of type 2 diabetes in two prospective cohorts of Chinese adults. <i>International Journal of Epidemiology</i> , 2016, 45, 1507-1516.	0.9	64
191	Association of Major Depressive Episodes With Stroke Risk in a Prospective Study of 0.5 Million Chinese Adults. <i>Stroke</i> , 2016, 47, 2203-2208.	1.0	27
192	Genetic variants within microRNA binding site of RAD51B are associated with risk of cervical cancer in Chinese women. <i>Cancer Medicine</i> , 2016, 5, 2596-2601.	1.3	18
193	Telomere structure and maintenance gene variants and risk of five cancer types. <i>International Journal of Cancer</i> , 2016, 139, 2655-2670.	2.3	43
194	Association of microRNA polymorphisms with the risk of head and neck squamous cell carcinoma in a Chinese population: a case-control study. <i>Chinese Journal of Cancer</i> , 2016, 35, 77.	4.9	33
195	Mitochondria-related miR-151a-5p reduces cellular ATP production by targeting CYTB in asthenozoospermia. <i>Scientific Reports</i> , 2016, 5, 17743.	1.6	52
196	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	5.8	86
197	Y chromosome haplogroups based genome-wide association study pinpoints revelation for interactions on non-obstructive azoospermia. <i>Scientific Reports</i> , 2016, 6, 33363.	1.6	7
198	Genome-wide association study identifies 8p21.3 associated with persistent hepatitis B virus infection among Chinese. <i>Nature Communications</i> , 2016, 7, 11664.	5.8	54

#	ARTICLE	IF	CITATIONS
199	Risk assessment model for invasive breast cancer in Hong Kong women. <i>Medicine (United States)</i> , 2016, 95, e4515.	0.4	11
200	Expression quantitative trait loci in long non-coding RNA PAX8-AS1 are associated with decreased risk of cervical cancer. <i>Molecular Genetics and Genomics</i> , 2016, 291, 1743-1748.	1.0	34
201	Genetic variants in chromatin-remodeling pathway associated with lung cancer risk in a Chinese population. <i>Gene</i> , 2016, 587, 178-182.	1.0	9
202	Role of <i>ATG10</i> expression quantitative trait loci in non-small cell lung cancer survival. <i>International Journal of Cancer</i> , 2016, 139, 1564-1573.	2.3	55
203	Systematic identification of genes with a cancer-testis expression pattern in 19 cancer types. <i>Nature Communications</i> , 2016, 7, 10499.	5.8	124
204	Meta-analysis of genome-wide association studies identifies multiple lung cancer susceptibility loci in never-smoking Asian women. <i>Human Molecular Genetics</i> , 2016, 25, 620-629.	1.4	50
205	Analysis of human papillomavirus 16 variants and risk for cervical cancer in Chinese population. <i>Virology</i> , 2016, 488, 156-161.	1.1	46
206	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
207	Fine mapping the MHC region identified four independent variants modifying susceptibility to chronic hepatitis B in Han Chinese. <i>Human Molecular Genetics</i> , 2016, 25, 1225-1232.	1.4	33
208	Genomic Landscape Survey Identifies SRSF1 as a Key Oncodriver in Small Cell Lung Cancer. <i>PLoS Genetics</i> , 2016, 12, e1005895.	1.5	144
209	Genetic variants in regulatory regions of microRNAs are associated with lung cancer risk. <i>Oncotarget</i> , 2016, 7, 47966-47974.	0.8	28
210	Copy number gain of VCX, X-linked multi-copy gene, leads to cell proliferation and apoptosis during spermatogenesis. <i>Oncotarget</i> , 2016, 7, 78532-78540.	0.8	11
211	X chromosome-wide identification of SNVs in microRNA genes and non-obstructive azoospermia risk in Han Chinese population. <i>Oncotarget</i> , 2016, 7, 49122-49129.	0.8	7
212	The driving role of cancer-testis genes: a new perspective for cancer researches. <i>Translational Cancer Research</i> , 2016, 5, S598-S600.	0.4	0
213	Potentially functional polymorphisms in PAK 1 are associated with risk of lung cancer in a Chinese population. <i>Cancer Medicine</i> , 2015, 4, 1781-1787.	1.3	6
214	Prediction models and risk assessment for silicosis using a retrospective cohort study among workers exposed to silica in China. <i>Scientific Reports</i> , 2015, 5, 11059.	1.6	25
215	The eQTL-missense polymorphisms of APOBEC3H are associated with lung cancer risk in a Han Chinese population. <i>Scientific Reports</i> , 2015, 5, 14969.	1.6	15
216	Mitochondria-related miR-141-3p contributes to mitochondrial dysfunction in HFD-induced obesity by inhibiting PTEN. <i>Scientific Reports</i> , 2015, 5, 16262.	1.6	48

#	ARTICLE	IF	CITATIONS
217	Hepatitis B virus genotype, mutations, human leukocyte antigen polymorphisms and their interactions in hepatocellular carcinoma: a multi-centre case-control study. <i>Scientific Reports</i> , 2015, 5, 16489.	1.6	35
218	Evaluation of regulatory genetic variants in POU5F1 and risk of congenital heart disease in Han Chinese. <i>Scientific Reports</i> , 2015, 5, 15860.	1.6	5
219	Genome-wide Association Study on Platinum-induced Hepatotoxicity in Non-Small Cell Lung Cancer Patients. <i>Scientific Reports</i> , 2015, 5, 11556.	1.6	23
220	Exploring causal associations of alcohol with cardiovascular and metabolic risk factors in a Chinese population using Mendelian randomization analysis. <i>Scientific Reports</i> , 2015, 5, 14005.	1.6	38
221	Clinical outcome and expression of mutant P53, P16, and Smad4 in lung adenocarcinoma: a prospective study. <i>World Journal of Surgical Oncology</i> , 2015, 13, 128.	0.8	18
222	Genetic variants associated with longer telomere length are associated with increased lung cancer risk among never-smoking women in Asia: a report from the female lung cancer consortium in Asia. <i>International Journal of Cancer</i> , 2015, 137, 311-319.	2.3	72
223	Cumulative Effect and Predictive Value of Genetic Variants Associated with Type 2 Diabetes in Han Chinese: A Case-Control Study. <i>PLoS ONE</i> , 2015, 10, e0116537.	1.1	19
224	A Common Variant Of Ubiquinol-Cytochrome c Reductase Complex Is Associated with DDH. <i>PLoS ONE</i> , 2015, 10, e0120212.	1.1	32
225	Expression Quantitative Trait Loci for CARD8 Contributes to Risk of Two Infection-Related Cancers—Hepatocellular Carcinoma and Cervical Cancer. <i>PLoS ONE</i> , 2015, 10, e0132352.	1.1	20
226	Low Goiter Rate Associated with Small Average Thyroid Volume in Schoolchildren after the Elimination of Iodine Deficiency Disorders. <i>PLoS ONE</i> , 2015, 10, e0141552.	1.1	17
227	Potentially Functional Polymorphisms in <i>POU5F1</i> Gene Are Associated with the Risk of Lung Cancer in Han Chinese. <i>BioMed Research International</i> , 2015, 2015, 1-7.	0.9	3
228	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
229	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
230	Genetic Variations in Key MicroRNAs are Associated With the Survival of Nonsmall Cell Lung Cancer. <i>Medicine (United States)</i> , 2015, 94, e2084.	0.4	14
231	Progress of cancer genomics. <i>Thoracic Cancer</i> , 2015, 6, 557-560.	0.8	1
232	Whole-exome sequencing identify a new mutation of MYH7 in a Chinese family with left ventricular noncompaction. <i>Gene</i> , 2015, 558, 138-142.	1.0	16
233	Systematical analyses of variants in CTCF-binding sites identified a novel lung cancer susceptibility locus among Chinese population. <i>Scientific Reports</i> , 2015, 5, 7833.	1.6	16
234	Interactions between household air pollution and GWAS-identified lung cancer susceptibility markers in the Female Lung Cancer Consortium in Asia (FLCCA). <i>Human Genetics</i> , 2015, 134, 333-341.	1.8	34

#	ARTICLE	IF	CITATIONS
235	Genome-wide Association Study of Survival in Early-stage Non-Small Cell Lung Cancer. <i>Annals of Surgical Oncology</i> , 2015, 22, 630-635.	0.7	52
236	Genetic variants of H2AX gene were associated with P M 2.5 -modulated DNA damage levels in Chinese Han populations. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2015, 778, 41-45.	0.4	3
237	A genome-wide association study of mitochondrial DNA in Chinese men identifies two risk single nucleotide substitutions for idiopathic oligoasthenospermia. <i>Mitochondrion</i> , 2015, 24, 87-92.	1.6	9
238	Joint effect of CENTD2 and KCNQ1 polymorphisms on the risk of type 2 diabetes mellitus among Chinese Han population. <i>Molecular and Cellular Endocrinology</i> , 2015, 407, 46-51.	1.6	12
239	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
240	Plasma mi<scp>RNA</scp>s as early biomarkers for detecting hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2015, 137, 1679-1690.	2.3	188
241	Low-Frequency Coding Variants at 6p21.33 and 20q11.21 Are Associated with Lung Cancer Risk in Chinese Populations. <i>American Journal of Human Genetics</i> , 2015, 96, 832-840.	2.6	41
242	Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. <i>Carcinogenesis</i> , 2015, 36, 1314-1326.	1.3	15
243	Associated Links Among Smoking, Chronic Obstructive Pulmonary Disease, and Small Cell Lung Cancer: A Pooled Analysis in the International Lung Cancer Consortium. <i>EBioMedicine</i> , 2015, 2, 1677-1685.	2.7	49
244	Oral lesions, chronic diseases and the risk of head and neck cancer. <i>Oral Oncology</i> , 2015, 51, 1082-1087.	0.8	31
245	Genetic variations in DROSHA and DICER and survival of advanced non-small cell lung cancer: a two-stage study in Chinese population. <i>Genes and Genomics</i> , 2015, 37, 571-578.	0.5	0
246	Association analysis identifies new risk loci for congenital heart disease in Chinese populations. <i>Nature Communications</i> , 2015, 6, 8082.	5.8	26
247	Polymorphisms in alternative splicing associated genes are associated with lung cancer risk in a Chinese population. <i>Lung Cancer</i> , 2015, 89, 238-242.	0.9	9
248	Telomere length, genetic variants and gastric cancer risk in a Chinese population. <i>Carcinogenesis</i> , 2015, 36, 963-970.	1.3	46
249	U-shaped association between telomere length and esophageal squamous cell carcinoma risk: a case-control study in Chinese population. <i>Frontiers of Medicine</i> , 2015, 9, 478-486.	1.5	10
250	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
251	Potentially functional polymorphisms in aminoacyl-tRNA synthetases genes are associated with breast cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2015, 54, 577-583.	1.3	15
252	Genome-wide association study in Chinese identifies novel loci for blood pressure and hypertension. <i>Human Molecular Genetics</i> , 2015, 24, 865-874.	1.4	157

#	ARTICLE	IF	CITATIONS
253	Pathway analysis for a genome-wide association study of pneumoconiosis. <i>Toxicology Letters</i> , 2015, 232, 284-292.	0.4	8
254	Circulating Tumor Cells Enriched by the Depletion of Leukocytes with Bi-Antibodies in Non-Small Cell Lung Cancer: Potential Clinical Application. <i>PLoS ONE</i> , 2015, 10, e0137076.	1.1	18
255	Genetic Variation in the 3'-Untranslated Region of NBN Gene Is Associated with Gastric Cancer Risk in a Chinese Population. <i>PLoS ONE</i> , 2015, 10, e0139059.	1.1	7
256	Genome-wide association study identifies two new susceptibility loci for colorectal cancer at 5q23.3 and 17q12 in Han Chinese. <i>Oncotarget</i> , 2015, 6, 40327-40336.	0.8	21
257	A functional variant in miR-155 regulation region contributes to lung cancer risk and survival. <i>Oncotarget</i> , 2015, 6, 42781-42792.	0.8	47
258	Family History of Cancer and Head and Neck Cancer Risk in a Chinese Population. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015, 16, 8003-8008.	0.5	5
259	Comparison of dimension reduction-based logistic regression models for case-control genome-wide association study: principal components analysis vs. partial least squares. <i>Journal of Biomedical Research</i> , 2015, 29, 298.	0.7	9
260	Spatio-Temporal Trends and Risk Factors for Shigella from 2001 to 2011 in Jiangsu Province, People's Republic of China. <i>PLoS ONE</i> , 2014, 9, e83487.	1.1	31
261	Genetic Variations in the Flanking Regions of miR-101-2 Are Associated with Increased Risk of Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e86319.	1.1	19
262	Genetic Variants at 10p11 Confer Risk of Tetralogy of Fallot in Chinese of Nanjing. <i>PLoS ONE</i> , 2014, 9, e89636.	1.1	6
263	Evaluation of the Impact of Hepatitis B Vaccination in Adults in Jiangsu Province, China. <i>PLoS ONE</i> , 2014, 9, e101501.	1.1	16
264	Association of GWAS-Identified Lung Cancer Susceptibility Loci with Survival Length in Patients with Small-Cell Lung Cancer Treated with Platinum-Based Chemotherapy. <i>PLoS ONE</i> , 2014, 9, e113574.	1.1	8
265	A genome-wide association study identifies susceptibility loci of silica-related pneumoconiosis in Han Chinese. <i>Human Molecular Genetics</i> , 2014, 23, 6385-6394.	1.4	24
266	A genome-wide gene-gene interaction analysis identifies an epistatic gene pair for lung cancer susceptibility in Han Chinese. <i>Carcinogenesis</i> , 2014, 35, 572-577.	1.3	29
267	Genetic variants at 8q24 are associated with risk of esophageal squamous cell carcinoma in a Chinese population. <i>Cancer Science</i> , 2014, 105, 731-735.	1.7	23
268	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
269	<i>TGFBR1</i> tagging SNPs and gastric cancer susceptibility: A two-stage case-control study in Chinese population. <i>Molecular Carcinogenesis</i> , 2014, 53, 109-116.	1.3	10
270	Association of polymorphisms at <i>HORMAD2</i> and prognosis in advanced non-small-cell lung cancer patients. <i>Cancer Epidemiology</i> , 2014, 38, 414-418.	0.8	5

#	ARTICLE	IF	CITATIONS
271	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
272	A genetic variant at KIF1B predicts clinical outcome of HBV-related hepatocellular carcinoma in Chinese. <i>Cancer Epidemiology</i> , 2014, 38, 608-612.	0.8	5
273	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
274	Genome-wide association study identifies three susceptibility loci for laryngeal squamous cell carcinoma in the Chinese population. <i>Nature Genetics</i> , 2014, 46, 1110-1114.	9.4	57
275	Genome-wide association study identifies new susceptibility loci for epithelial ovarian cancer in Han Chinese women. <i>Nature Communications</i> , 2014, 5, 4682.	5.8	59
276	Polymorphisms in MicroRNAs Are Associated with Survival in Non-Small Cell Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2503-2511.	1.1	22
277	A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. <i>Carcinogenesis</i> , 2014, 35, 1528-1535.	1.3	47
278	Evaluation of functional genetic variants at 6q25.1 and risk of breast cancer in a Chinese population. <i>Breast Cancer Research</i> , 2014, 16, 422.	2.2	21
279	Genetic variants in SMARCA4 genes are associated with DNA damage levels in Chinese population. <i>Toxicology Letters</i> , 2014, 229, 327-332.	0.4	6
280	Association analysis identifies new risk loci for non-obstructive azoospermia in Chinese men. <i>Nature Communications</i> , 2014, 5, 3857.	5.8	64
281	Gene copy number alterations in the azoospermia-associated AZFc region and their effect on spermatogenic impairment. <i>Molecular Human Reproduction</i> , 2014, 20, 836-843.	1.3	27
282	The Spatial Analysis on Hemorrhagic Fever with Renal Syndrome in Jiangsu Province, China Based on Geographic Information System. <i>PLoS ONE</i> , 2014, 9, e83848.	1.1	14
283	A Genetic Variant in the Promoter Region of miR-106b-25 Cluster Predict Clinical Outcome of HBV-Related Hepatocellular Carcinoma in Chinese. <i>PLoS ONE</i> , 2014, 9, e85394.	1.1	23
284	Population Aging and Migrant Workers: Bottlenecks in Tuberculosis Control in Rural China. <i>PLoS ONE</i> , 2014, 9, e88290.	1.1	28
285	A Genetic Variant in Primary miR-378 Is Associated with Risk and Prognosis of Hepatocellular Carcinoma in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e93707.	1.1	24
286	Replication of the 4p16 Susceptibility Locus in Congenital Heart Disease in Han Chinese Populations. <i>PLoS ONE</i> , 2014, 9, e107411.	1.1	14
287	A genetic variant in pseudogene E2F3P1 contributes to prognosis of hepatocellular carcinoma. <i>Journal of Biomedical Research</i> , 2014, 28, 194.	0.7	12
288	Genetic variants at 10q23.33 are associated with plasma lipid levels in a Chinese population. <i>Journal of Biomedical Research</i> , 2014, 28, 53-8.	0.7	5

#	ARTICLE	IF	CITATIONS
289	Genetic variants at 4q21, 4q23 and 12q24 are associated with esophageal squamous cell carcinoma risk in a Chinese population. <i>Human Genetics</i> , 2013, 132, 649-656.	1.8	34
290	New loci associated with chronic hepatitis B virus infection in Han Chinese. <i>Nature Genetics</i> , 2013, 45, 1499-1503.	9.4	140
291	Genome-wide microRNA expression profiling in idiopathic non-obstructive azoospermia: significant up-regulation of miR-141, miR-429 and miR-7-1-3p. <i>Human Reproduction</i> , 2013, 28, 1827-1836.	0.4	115
292	Genetic variant in fat mass and obesity-associated gene associated with type 2 diabetes risk in Han Chinese. <i>BMC Genetics</i> , 2013, 14, 86.	2.7	16
293	IL-23R polymorphisms, HBV infection, and risk of hepatocellular carcinoma in a high-risk Chinese population. <i>Journal of Gastroenterology</i> , 2013, 48, 125-131.	2.3	30
294	Potentially functional polymorphisms in ATG10 are associated with risk of breast cancer in a Chinese population. <i>Gene</i> , 2013, 527, 491-495.	1.0	44
295	Genetic variants at 10q23 are associated with risk of head and neck cancer in a Chinese population. <i>Oral Oncology</i> , 2013, 49, 332-335.	0.8	12
296	Evaluation of genetic variants in microRNA biosynthesis genes and risk of breast cancer in Chinese women. <i>International Journal of Cancer</i> , 2013, 133, 2216-2224.	2.3	50
297	Genome-wide association study identifies common variants in SLC39A6 associated with length of survival in esophageal squamous-cell carcinoma. <i>Nature Genetics</i> , 2013, 45, 632-638.	9.4	97
298	Genetic variants in STAT4 and HLA-DQ genes confer risk of hepatitis B virus-related hepatocellular carcinoma. <i>Nature Genetics</i> , 2013, 45, 72-75.	9.4	259
299	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
300	Genetic variants in HLA-DP/DQ contribute to risk of cervical cancer: A two-stage study in Chinese women. <i>Gynecologic Oncology</i> , 2013, 129, 401-405.	0.6	19
301	A genome-wide association study identifies two risk loci for congenital heart malformations in Han Chinese populations. <i>Nature Genetics</i> , 2013, 45, 818-821.	9.4	88
302	Genome-Wide Association Study Identifies a Novel Susceptibility Locus at 12q23.1 for Lung Squamous Cell Carcinoma in Han Chinese. <i>PLoS Genetics</i> , 2013, 9, e1003190.	1.5	41
303	Genetic variation in a hsa-let-7 binding site in RAD52 is associated with breast cancer susceptibility. <i>Carcinogenesis</i> , 2013, 34, 689-693.	1.3	43
304	DAZ duplications confer the predisposition of Y chromosome haplogroup K* to non-obstructive azoospermia in Han Chinese populations. <i>Human Reproduction</i> , 2013, 28, 2440-2449.	0.4	13
305	Genetic variants at 5p15 are associated with risk and early onset of gastric cancer in Chinese populations. <i>Carcinogenesis</i> , 2013, 34, 2539-2542.	1.3	13
306	Nonsynonymous polymorphisms in <i>FAT4</i> gene are associated with the risk of esophageal cancer in an Eastern Chinese population. <i>International Journal of Cancer</i> , 2013, 133, 357-361.	2.3	10

#	ARTICLE	IF	CITATIONS
307	Human genome epidemiology, progress and future. <i>Journal of Biomedical Research</i> , 2013, 27, 167-169.	0.7	17
308	Prognostic assessment of apoptotic gene polymorphisms in non-small cell lung cancer in Chinese. <i>Journal of Biomedical Research</i> , 2013, 27, 231-8.	0.7	6
309	Genome-wide analysis of runs of homozygosity identifies new susceptibility regions of lung cancer in Han Chinese. <i>Journal of Biomedical Research</i> , 2013, 27, 208.	0.7	18
310	GWAS Identifies Novel Susceptibility Loci on 6p21.32 and 21q21.3 for Hepatocellular Carcinoma in Chronic Hepatitis B Virus Carriers. <i>PLoS Genetics</i> , 2012, 8, e1002791.	1.5	177
311	Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , 2012, 21, 4980-4995.	1.4	196
312	Class A Scavenger Receptor Deficiency Exacerbates Lung Tumorigenesis by Cultivating a Procarcinogenic Microenvironment in Humans and Mice. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2012, 186, 763-772.	2.5	23
313	Genetic variants on chromosome 6p21.1 and 6p22.3 are associated with type 2 diabetes risk: a case-control study in Han Chinese. <i>Journal of Human Genetics</i> , 2012, 57, 320-325.	1.1	25
314	Genetic Variants at 14q24.1 and Breast Cancer Susceptibility: a Fine-Mapping Study in Chinese Women. <i>DNA and Cell Biology</i> , 2012, 31, 1114-1120.	0.9	4
315	Association analyses identify multiple new lung cancer susceptibility loci and their interactions with smoking in the Chinese population. <i>Nature Genetics</i> , 2012, 44, 895-899.	9.4	129
316	Serum microRNA profiling and breast cancer risk: the use of miR-484/191 as endogenous controls. <i>Carcinogenesis</i> , 2012, 33, 828-834.	1.3	193
317	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
318	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335.	9.4	286
319	Breast cancer risk assessment with five independent genetic variants and two risk factors in Chinese women. <i>Breast Cancer Research</i> , 2012, 14, R17.	2.2	57
320	Common Variant rs1017 Is Not Associated with Susceptibility to Congenital Heart Disease in a Chinese Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2012, 16, 679-683.	0.3	9
321	Genome-Wide Association Study of Prognosis in Advanced Non-Small Cell Lung Cancer Patients Receiving Platinum-Based Chemotherapy. <i>Clinical Cancer Research</i> , 2012, 18, 5507-5514.	3.2	56
322	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
323	A germline variant N375S in MET and gastric cancer susceptibility in a Chinese population. <i>Journal of Biomedical Research</i> , 2012, 26, 315-318.	0.7	11
324	Genome-wide association study in Chinese men identifies two new prostate cancer risk loci at 9q31.2 and 19q13.4. <i>Nature Genetics</i> , 2012, 44, 1231-1235.	9.4	160

#	ARTICLE	IF	CITATIONS
325	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
326	A Genetic Variant in 3' Untranslated Region of Cyclooxygenases-2 Gene Is Associated with Risk of Gastric Cancer in a Chinese Population. <i>DNA and Cell Biology</i> , 2012, 31, 1252-1257.	0.9	6
327	A genome-wide association study in Chinese men identifies three risk loci for non-obstructive azoospermia. <i>Nature Genetics</i> , 2012, 44, 183-186.	9.4	139
328	A Genetic Variant in the Promoter Region of miR-106b-25 Cluster and Risk of HBV Infection and Hepatocellular Carcinoma. <i>PLoS ONE</i> , 2012, 7, e32230.	1.1	69
329	Genetic Variants of IDE-KIF11-HHEX at 10q23.33 Associated with Type 2 Diabetes Risk: A Fine-Mapping Study in Chinese Population. <i>PLoS ONE</i> , 2012, 7, e35060.	1.1	32
330	Potentially functional polymorphisms in DNA repair genes and non-small cell lung cancer survival: A pathway-based analysis. <i>Molecular Carcinogenesis</i> , 2012, 51, 546-552.	1.3	48
331	Common genetic variants in <i>TERT</i> contribute to risk of cervical cancer in a Chinese population. <i>Molecular Carcinogenesis</i> , 2012, 51, E118-22.	1.3	28
332	Smoking and Genetic Risk Variation Across Populations of European, Asian, and African Ancestry: A Meta-Analysis of Chromosome 15q25. <i>Genetic Epidemiology</i> , 2012, 36, 340-351.	0.6	69
333	Genetic variants in human leukocyte antigen/DP-DQ influence both hepatitis B virus clearance and hepatocellular carcinoma development. <i>Hepatology</i> , 2012, 55, 1426-1431.	3.6	157
334	Host immune gene polymorphisms were associated with the prognosis of non-small cell lung cancer in Chinese. <i>International Journal of Cancer</i> , 2012, 130, 671-676.	2.3	13
335	Potentially functional polymorphisms in IL23 receptor and risk of esophageal cancer in a Chinese population. <i>International Journal of Cancer</i> , 2012, 130, 1093-1097.	2.3	32
336	Identification of ten serum microRNAs from a genome-wide serum microRNA expression profile as novel noninvasive biomarkers for nonsmall cell lung cancer diagnosis. <i>International Journal of Cancer</i> , 2012, 130, 1620-1628.	2.3	251
337	A Genetic Variant in Long Non-Coding RNA HULC Contributes to Risk of HBV-Related Hepatocellular Carcinoma in a Chinese Population. <i>PLoS ONE</i> , 2012, 7, e35145.	1.1	145
338	A genome-wide association study identifies two new lung cancer susceptibility loci at 13q12.12 and 22q12.2 in Han Chinese. <i>Nature Genetics</i> , 2011, 43, 792-796.	9.4	340
339	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
340	Potentially functional polymorphisms in cell cycle genes and the survival of non-small cell lung cancer in a Chinese population. <i>Lung Cancer</i> , 2011, 73, 32-37.	0.9	22
341	Polymorphisms of key chemokine genes and survival of non-small cell lung cancer in Chinese. <i>Lung Cancer</i> , 2011, 74, 164-169.	0.9	17
342	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

#	ARTICLE	IF	CITATIONS
343	Genetic susceptibility of cervical cancer. <i>Journal of Biomedical Research</i> , 2011, 25, 155-164.	0.7	34
344	A variant affecting miRNAs binding in the circadian gene Neuronal PAS domain protein 2 (NPAS2) is not associated with breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2011, 127, 769-775.	1.1	11
345	Genetic variants in ultraconserved elements and risk of breast cancer in Chinese population. <i>Breast Cancer Research and Treatment</i> , 2011, 128, 855-861.	1.1	11
346	Genetic variants of 6q25 and breast cancer susceptibility: a two-stage fine mapping study in a Chinese population. <i>Breast Cancer Research and Treatment</i> , 2011, 129, 901-907.	1.1	18
347	<i>MDM2</i> SNP309 contributes to non-small cell lung cancer survival in Chinese. <i>Molecular Carcinogenesis</i> , 2011, 50, 433-438.	1.3	19
348	A potentially functional polymorphism in the promoter region of miR-34b/c is associated with an increased risk for primary hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2011, 128, 412-417.	2.3	169
349	Genetic Polymorphisms in the Precursor MicroRNA Flanking Region and Non-Small Cell Lung Cancer Survival. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2011, 183, 641-648.	2.5	46
350	Replication and Functional Genomic Analyses of the Breast Cancer Susceptibility Locus at 6q25.1 Generalize Its Importance in Women of Chinese, Japanese, and European Ancestry. <i>Cancer Research</i> , 2011, 71, 1344-1355.	0.4	71
351	Variations in <i>HSPA1B</i> at 6p21.3 Are Associated with Lung Cancer Risk and Prognosis in Chinese Populations. <i>Cancer Research</i> , 2011, 71, 7576-7586.	0.4	30
352	Genetic variants at 1q22 and 10q23 reproducibly associated with gastric cancer susceptibility in a Chinese population. <i>Carcinogenesis</i> , 2011, 32, 848-852.	1.3	73
353	Genetic Variants at 1p11.2 and Breast Cancer Risk: A Two-Stage Study in Chinese Women. <i>PLoS ONE</i> , 2011, 6, e21563.	1.1	24
354	Prognostic Significance of Survivin Polymorphisms on Non-small Cell Lung Cancer Survival. <i>Journal of Thoracic Oncology</i> , 2010, 5, 1748-1754.	0.5	26
355	Circulating IL-1 β levels, polymorphisms of IL-1B, and risk of cervical cancer in Chinese women. <i>Journal of Cancer Research and Clinical Oncology</i> , 2010, 136, 709-716.	1.2	62
356	Potentially functional polymorphisms in ESR1 and breast cancer risk: a meta-analysis. <i>Breast Cancer Research and Treatment</i> , 2010, 121, 177-184.	1.1	49
357	Genetic variants in trinucleotide repeat-containing 9 (TNRC9) are associated with risk of estrogen receptor positive breast cancer in a Chinese population. <i>Breast Cancer Research and Treatment</i> , 2010, 124, 237-241.	1.1	28
358	Polymorphisms involved in the miR-218-LAMB3 pathway and susceptibility of cervical cancer, a case-control study in Chinese women. <i>Gynecologic Oncology</i> , 2010, 117, 287-290.	0.6	64
359	Genetic variants in one-carbon metabolism-related genes contribute to NSCLC prognosis in a Chinese population. <i>Cancer</i> , 2010, 116, 5700-5709.	2.0	14
360	Genetic variation of <i>PSCA</i> gene is associated with the risk of both diffuse and intestinal-type gastric cancer in a Chinese population. <i>International Journal of Cancer</i> , 2010, 127, 2183-2189.	2.3	72

#	ARTICLE	IF	CITATIONS
361	A nonsynonymous polymorphism in <i>IL23R</i> gene is associated with risk of gastric cancer in a Chinese population. <i>Molecular Carcinogenesis</i> , 2010, 49, 862-868.	1.3	51
362	Genome-wide association study identifies 1p36.22 as a new susceptibility locus for hepatocellular carcinoma in chronic hepatitis B virus carriers. <i>Nature Genetics</i> , 2010, 42, 755-758.	9.4	319
363	A variant in the CHEK2 promoter at a methylation site relieves transcriptional repression and confers reduced risk of lung cancer. <i>Carcinogenesis</i> , 2010, 31, 1251-1258.	1.3	26
364	International Lung Cancer Consortium: Coordinated association study of 10 potential lung cancer susceptibility variants. <i>Carcinogenesis</i> , 2010, 31, 625-633.	1.3	56
365	MTHFR c.1793G>A polymorphism is associated with congenital cardiac disease in a Chinese population. <i>Cardiology in the Young</i> , 2010, 20, 318-326.	0.4	20
366	Reply to H.M. Heneghan et al. <i>Journal of Clinical Oncology</i> , 2010, 28, e575-e576.	0.8	3
367	The 5p15.33 Locus Is Associated with Risk of Lung Adenocarcinoma in Never-Smoking Females in Asia. <i>PLoS Genetics</i> , 2010, 6, e1001051.	1.5	168
368	A 3'-Untranslated Region Polymorphism in IGF1 Predicts Survival of Non-Small Cell Lung Cancer in a Chinese Population. <i>Clinical Cancer Research</i> , 2010, 16, 1236-1244.	3.2	41
369	Polymorphisms in EGFR and VEGF contribute to non-small-cell lung cancer survival in a Chinese population. <i>Carcinogenesis</i> , 2010, 31, 1080-1086.	1.3	36
370	Serum MicroRNA Signatures Identified in a Genome-Wide Serum MicroRNA Expression Profiling Predict Survival of Non-Small-Cell Lung Cancer. <i>Journal of Clinical Oncology</i> , 2010, 28, 1721-1726.	0.8	759
371	Candidate variants at 6p21.33 and 6p22.1 and risk of non-small cell lung cancer in a Chinese population. <i>International Journal of Molecular Epidemiology and Genetics</i> , 2010, 1, 11-8.	0.4	8
372	Functional polymorphisms in two pre-microRNAs and cancer risk: a meta-analysis. <i>International Journal of Molecular Epidemiology and Genetics</i> , 2010, 1, 358-66.	0.4	21
373	Cigarette Smoking, MDM2SNP309, Gene-Environment Interactions, and Lung Cancer Risk: A Meta-Analysis. <i>Journal of Toxicology and Environmental Health - Part A: Current Issues</i> , 2009, 72, 677-682.	1.1	24
374	Genetic Polymorphisms in <i>Cytotoxic T-Lymphocyte Antigen 4</i> and Cancer: The Dialectical Nature of Subtle Human Immune Dysregulation. <i>Cancer Research</i> , 2009, 69, 6011-6014.	0.4	58
375	Common genetic variants on 5p15.33 contribute to risk of lung adenocarcinoma in a Chinese population. <i>Carcinogenesis</i> , 2009, 30, 987-990.	1.3	72
376	Functional characterization of a promoter polymorphism in <i>APE1/Ref-1</i> that contributes to reduced lung cancer susceptibility. <i>FASEB Journal</i> , 2009, 23, 3459-3469.	0.2	65
377	Genetic Variants on Chromosome 15q25 Associated with Lung Cancer Risk in Chinese Populations. <i>Cancer Research</i> , 2009, 69, 5065-5072.	0.4	138
378	Association of human aryl hydrocarbon receptor gene polymorphisms with risk of lung cancer among cigarette smokers in a Chinese population. <i>Pharmacogenetics and Genomics</i> , 2009, 19, 25-34.	0.7	49

#	ARTICLE	IF	CITATIONS
379	A Functional Genetic Variant in microRNA-196a2 Is Associated with Increased Susceptibility of Lung Cancer in Chinese. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1183-1187.	1.1	278
380	Polymorphisms in HPV E6/E7 protein interacted genes and risk of cervical cancer in Chinese women: A case-control analysis. <i>Gynecologic Oncology</i> , 2009, 114, 327-331.	0.6	28
381	Common genetic variants in pre-microRNAs were associated with increased risk of breast cancer in Chinese women. <i>Human Mutation</i> , 2009, 30, 79-84.	1.1	356
382	Functional variant in microRNA-196a2 contributes to the susceptibility of congenital heart disease in a Chinese population. <i>Human Mutation</i> , 2009, 30, 1231-1236.	1.1	124
383	Putative functional polymorphisms of <i>MMP9</i> predict survival of NSCLC in a Chinese population. <i>International Journal of Cancer</i> , 2009, 124, 2172-2178.	2.3	22
384	Genetic susceptibility of lung cancer associated with common variants in the 3' untranslated regions of the adenosine triphosphate-binding cassette B1 (<i>ABCB1</i>) and <i>ABCC1</i> candidate transporter genes for carcinogen export. <i>Cancer</i> , 2009, 115, 595-607.	2.0	30
385	Association between polymorphisms in the <i>GSTA4</i> gene and risk of lung cancer: A case-control study in a Southeastern Chinese population. <i>Molecular Carcinogenesis</i> , 2009, 48, 253-259.	1.3	7
386	Polymorphisms in BRCA1, BRCA1-interacting genes and susceptibility of breast cancer in Chinese women. <i>Journal of Cancer Research and Clinical Oncology</i> , 2009, 135, 1569-1575.	1.2	29
387	CCND1 G870A polymorphism contributes to breast cancer susceptibility: a meta-analysis. <i>Breast Cancer Research and Treatment</i> , 2009, 116, 571-575.	1.1	30
388	A polymorphism in Werner syndrome gene is associated with breast cancer susceptibility in Chinese women. <i>Breast Cancer Research and Treatment</i> , 2009, 118, 169-175.	1.1	43
389	ERCC6/CSB gene polymorphisms and lung cancer risk. <i>Cancer Letters</i> , 2009, 273, 172-176.	3.2	26
390	Genetic variants in GTF2H1 and risk of lung cancer: A case-control analysis in a Chinese population. <i>Lung Cancer</i> , 2009, 63, 180-186.	0.9	10
391	EGF promoter SNPs, plasma EGF levels and risk of breast cancer in Chinese women. <i>Breast Cancer Research and Treatment</i> , 2008, 111, 321-327.	1.1	21
392	TGFB1 and TGFBR2 functional polymorphisms and risk of esophageal squamous cell carcinoma: a case-control analysis in a Chinese population. <i>Journal of Cancer Research and Clinical Oncology</i> , 2008, 134, 345-351.	1.2	40
393	Polymorphisms in hMLH1 and risk of early-onset lung cancer in a southeast Chinese population. <i>Lung Cancer</i> , 2008, 59, 164-170.	0.9	30
394	Potentially functional polymorphisms of EXO1 and risk of lung cancer in a Chinese population: A case-control analysis. <i>Lung Cancer</i> , 2008, 60, 340-346.	0.9	37
395	Tagging single nucleotide polymorphisms in MBD4 are associated with risk of lung cancer in a Chinese population. <i>Lung Cancer</i> , 2008, 62, 281-286.	0.9	15
396	Association of Common <i>PALB2</i> Polymorphisms with Breast Cancer Risk: A Case-Control Study. <i>Clinical Cancer Research</i> , 2008, 14, 5931-5937.	3.2	38

#	ARTICLE	IF	CITATIONS
397	Genetic variants in fibroblast growth factor receptor 2 (FGFR2) contribute to susceptibility of breast cancer in Chinese women. <i>Carcinogenesis</i> , 2008, 29, 2341-2346.	1.3	88
398	A tandem repeat of human telomerase reverse transcriptase (hTERT) and risk of breast cancer development and metastasis in Chinese women. <i>Carcinogenesis</i> , 2008, 29, 1197-1201.	1.3	34
399	Functional Genetic Variations in <i>Cytotoxic T-Lymphocyte Antigen 4</i> and Susceptibility to Multiple Types of Cancer. <i>Cancer Research</i> , 2008, 68, 7025-7034.	0.4	151
400	Tagging Single Nucleotide Polymorphisms in Phosphoinositide-3-Kinase-Related Protein Kinase Genes Involved in DNA Damage Checkpoints and Lung Cancer Susceptibility. <i>Clinical Cancer Research</i> , 2008, 14, 2887-2891.	3.2	9
401	International Lung Cancer Consortium: Pooled Analysis of Sequence Variants in DNA Repair and Cell Cycle Pathways. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3081-3089.	1.1	93
402	Genetic variants in peroxisome proliferator-activated receptor- β gene are associated with risk of lung cancer in a Chinese population. <i>Carcinogenesis</i> , 2008, 29, 342-350.	1.3	24
403	Methyl-CpG binding domain 1 gene polymorphisms and lung cancer risk in a Chinese population. <i>Biomarkers</i> , 2008, 13, 607-617.	0.9	6
404	Genetic variants of miRNA sequences and non-small cell lung cancer survival. <i>Journal of Clinical Investigation</i> , 2008, 118, 2600-8.	3.9	485
405	MDM2 Promoter Polymorphism SNP309 Contributes to Tumor Susceptibility: Evidence from 21 Case-Control Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 2717-2723.	1.1	100
406	Genetic variants in MGMT and risk of lung cancer in Southeastern Chinese: a haplotype-based analysis. <i>Human Mutation</i> , 2007, 28, 431-440.	1.1	46
407	Variant alleles of TGFB1 and TGFB2 are associated with a decreased risk of gastric cancer in a Chinese population. <i>International Journal of Cancer</i> , 2007, 120, 1330-1335.	2.3	70
408	Variant genotypes and haplotypes of the epidermal growth factor gene promoter are associated with a decreased risk of gastric cancer in a high-risk Chinese population. <i>Cancer Science</i> , 2007, 98, 864-868.	1.7	35
409	Sequence variations in DNA repair gene XPC is associated with lung cancer risk in a Chinese population: a case-control study. <i>BMC Cancer</i> , 2007, 7, 81.	1.1	31
410	Common non-synonymous polymorphisms in the BRCA1 Associated RING Domain (BARD1) gene are associated with breast cancer susceptibility: a case-control analysis. <i>Breast Cancer Research and Treatment</i> , 2007, 102, 329-337.	1.1	25
411	The association of Y chromosome haplogroups with spermatogenic failure in the Han Chinese. <i>Journal of Human Genetics</i> , 2007, 52, 659-663.	1.1	12
412	Joint effects of single nucleotide polymorphisms in P53BP1 and p53 on breast cancer risk in a Chinese population. <i>Carcinogenesis</i> , 2006, 27, 766-771.	1.3	39
413	Allele 2 of the interleukin-1 receptor antagonist gene (IL1RN*2) is associated with a decreased risk of primary lung cancer. <i>Cancer Letters</i> , 2006, 236, 269-275.	3.2	41
414	Polymorphisms in the MDM2 promoter and risk of breast cancer: a case-control analysis in a Chinese population. <i>Cancer Letters</i> , 2006, 240, 261-267.	3.2	67

#	ARTICLE	IF	CITATIONS
415	Reduced folate carrier gene G80A polymorphism is associated with an increased risk of gastroesophageal cancers in a chinese population. <i>European Journal of Cancer</i> , 2006, 42, 3206-3211.	1.3	29
416	Genetic variants in the MDM2 promoter and lung cancer risk in a Chinese population. <i>International Journal of Cancer</i> , 2006, 118, 1275-1278.	2.3	66
417	Functional variants in the promoter of interleukin-1 β are associated with an increased risk of breast cancer: A case-control analysis in a Chinese population. <i>International Journal of Cancer</i> , 2006, 118, 2554-2558.	2.3	37
418	Variant genotypes of CDKN1A and CDKN1B are associated with an increased risk of breast cancer in Chinese women. <i>International Journal of Cancer</i> , 2006, 119, 2173-2178.	2.3	41
419	Polymorphisms in DNA damage binding protein 2 (DDB2) and susceptibility of primary lung cancer in the Chinese: a case-control study. <i>Carcinogenesis</i> , 2006, 27, 1475-1480.	1.3	20
420	A promoter polymorphism (A>T) of DNA repair gene XRCC1 is associated with risk of lung cancer in relation to tobacco smoking. <i>Pharmacogenetics and Genomics</i> , 2005, 15, 457-463.	0.7	55
421	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
422	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
423	Functional Polymorphisms of Matrix Metalloproteinase-9 Are Associated with Risk of Occurrence and Metastasis of Lung Cancer. <i>Clinical Cancer Research</i> , 2005, 11, 5433-5439.	3.2	96
424	Methylenetetrahydrofolate reductase polymorphisms/haplotypes and risk of gastric cancer: a case-control analysis in China. <i>Oncology Reports</i> , 2005, 13, 355-60.	1.2	40
425	Polymorphisms of DNA repair gene XRCC3 Thr241Met and risk of gastric cancer in a Chinese population. <i>Cancer Letters</i> , 2004, 206, 51-58.	3.2	44
426	Interleukin-1B gene promoter variants are associated with an increased risk of gastric cancer in a Chinese population. <i>Cancer Letters</i> , 2004, 215, 191-198.	3.2	84
427	P53 codon 72 polymorphism and risk of gastric cancer in a Chinese population. <i>Oncology Reports</i> , 2004, 11, 1115-20.	1.2	46
428	Smoking, DNA repair capacity and risk of nonsmall cell lung cancer. <i>International Journal of Cancer</i> , 2003, 107, 84-88.	2.3	125
429	p53 Codon 72 Arg Homozygotes Are Associated with an Increased Risk of Cutaneous Melanoma. <i>Journal of Investigative Dermatology</i> , 2003, 121, 1510-1514.	0.3	45
430	Dietary folate intake and lung cancer risk in former smokers: a case-control analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2003, 12, 980-6.	1.1	24
431	Modulation of repair of ultraviolet damage in the host-cell reactivation assay by polymorphic XPC and XPD/ERCC2 genotypes. <i>Carcinogenesis</i> , 2002, 23, 295-299.	1.3	248
432	Polymorphism of DNA ligase I and risk of lung cancer—a case-control analysis. <i>Lung Cancer</i> , 2002, 36, 243-247.	0.9	17

#	ARTICLE	IF	CITATIONS
433	P53 codon 72 polymorphism and risk of squamous cell carcinoma of the head and neck: a case-control study. <i>Cancer Letters</i> , 2002, 183, 123-130.	3.2	94
434	A variant of the DNA repair gene XRCC3 and risk of squamous cell carcinoma of the head and neck: A case-control analysis. <i>International Journal of Cancer</i> , 2002, 99, 869-872.	2.3	52
435	A novel polymorphism in human cytosine DNA-methyltransferase-3B promoter is associated with an increased risk of lung cancer. <i>Cancer Research</i> , 2002, 62, 4992-5.	0.4	137
436	Polymorphisms of 5,10-methylenetetrahydrofolate reductase and risk of gastric cancer in a Chinese population: A case-control study. <i>International Journal of Cancer</i> , 2001, 95, 332-336.	2.3	119
437	Cyclin D1 polymorphism and risk for squamous cell carcinoma of the head and neck: a case-control study. <i>Carcinogenesis</i> , 2001, 22, 1195-1199.	1.3	109
438	Polymorphisms of the DNA repair gene XRCC1 and risk of gastric cancer in a Chinese population. <i>International Journal of Cancer</i> , 2000, 88, 601-606.	2.3	165
439	Association of current income and reduction in income during the COVID-19 pandemic with anxiety and depression among non-healthcare workers. <i>Journal of Mental Health</i> , 0, , 1-12.	1.0	5