## Xiaoping Miao

List of Publications by Year in descending order

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286 20,413 64 papers citations h-index

g-index
35393

293
all docs doc

293 docs citations 293 times ranked

35393 citing authors

129

#	Article	IF	Citations
1	Urinary bisphenol A and its interaction with CYP17A1 rs743572 are associated with breast cancer risk. Chemosphere, 2022, 286, 131880.	4.2	8
2	Genome-wide gene-bisphenol A, F and triclosan interaction analyses on urinary oxidative stress markers. Science of the Total Environment, 2022, 807, 150753.	3.9	11
3	No Evidence for a Causal Link between Serum Uric Acid and Nonalcoholic Fatty Liver Disease from the Dongfeng-Tongji Cohort Study. Oxidative Medicine and Cellular Longevity, 2022, 2022, 1-10.	1.9	5
4	Aberrant RNA Splicing Is a Primary Link between Genetic Variation and Pancreatic Cancer Risk. Cancer Research, 2022, 82, 2084-2096.	0.4	14
5	Systematic analysis on expression quantitative trait loci identifies a novel regulatory variant in ring finger and WD repeat domain 3 associated with prognosis of pancreatic cancer. Chinese Medical Journal, 2022, 135, 1348-1357.	0.9	3
6	Potential Influence of Menstrual Status and Sex Hormones on Female Severe Acute Respiratory Syndrome Coronavirus 2 Infection: A Cross-sectional Multicenter Study in Wuhan, China. Clinical Infectious Diseases, 2021, 72, e240-e248.	2.9	96
7	CancerlmmunityQTL: a database to systematically evaluate the impact of genetic variants on immune infiltration in human cancer. Nucleic Acids Research, 2021, 49, D1065-D1073.	6.5	22
8	Repurposed Tocilizumab in Patients with Severe COVID-19. Journal of Immunology, 2021, 206, 599-606.	0.4	17
9	COVID-19 in Peritoneal Dialysis Patients. Clinical Journal of the American Society of Nephrology: CJASN, 2021, 16, 121-123.	2.2	44
10	Identification of genetic variants in m6A modification genes associated with pancreatic cancer risk in the Chinese population. Archives of Toxicology, 2021, 95, 1117-1128.	1.9	17
11	Analysis of Ovarian Injury Associated With COVID-19 Disease in Reproductive-Aged Women in Wuhan, China: An Observational Study. Frontiers in Medicine, 2021, 8, 635255.	1.2	86
12	Genetic variants associated with expression of TCF19 contribute to the risk of head and neck cancer in Chinese population. Journal of Medical Genetics, 2021, , jmedgenet-2020-107410.	1.5	7
13	Aberrant MCM10 SUMOylation induces genomic instability mediated by a genetic variant associated with survival of esophageal squamous cell carcinoma. Clinical and Translational Medicine, 2021, 11, e485.	1.7	8
14	Implications of Lifestyle Factors and Polygenic Risk Score for Absolute Risk Prediction of Colorectal Neoplasm and Risk-Adapted Screening. Frontiers in Molecular Biosciences, 2021, 8, 685410.	1.6	4
15	A genetic variant conferred high expression of CAV2 promotes pancreatic cancer progression and associates with poor prognosis. European Journal of Cancer, 2021, 151, 94-105.	1.3	10
16	Leveraging Fecal Microbial Markers to Improve the Diagnostic Accuracy of the Fecal Immunochemical Test for Advanced Colorectal Adenoma. Clinical and Translational Gastroenterology, 2021, 12, e00389.	1.3	7
17	Colorectal cancer risk variant rs7017386 modulates two oncogenic lncRNAs expression via ATF1-mediated long-range chromatin loop. Cancer Letters, 2021, 518, 140-151.	3.2	9
18	Bisphenol A exposure, interaction with genetic variants and colorectal cancer via mediating oxidative stress biomarkers. Environmental Pollution, 2021, 287, 117630.	3.7	11

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19	Associations of polycyclic aromatic hydrocarbons exposure and its interaction with XRCC1 genetic polymorphism with lung cancer: A case-control study. Environmental Pollution, 2021, 290, 118077.	3.7	6
20	Hepatocellular carcinoma risk variant modulates lncRNA HLA-DQB1-AS1 expression via a long-range enhancer–promoter interaction. Carcinogenesis, 2021, 42, 1347-1356.	1.3	10
21	FOXA1 of regulatory variant associated with risk of breast cancer through allele-specific enhancer in the Chinese population. Breast Cancer, $2021, 1.$	1.3	О
22	Association of blood pressure and longâ€term change with chronic kidney disease risk among Chinese adults with different glucose metabolism according to the 2017 ACC/AHA guidelines. Journal of Clinical Hypertension, 2021, , .	1.0	2
23	Prediction of 5-year risk of diabetes mellitus in relatively low risk middle-aged and elderly adults. Acta Diabetologica, 2020, 57, 63-70.	1.2	10
24	Urinary biomarkers of phthalates exposure and risks of thyroid cancer and benign nodule. Journal of Hazardous Materials, 2020, 383, 121189.	6.5	46
25	<i>ANKLE1</i> N <sup>6</sup> â€Methyladenosineâ€related variant is associated with colorectal cancer risk by maintaining the genomic stability. International Journal of Cancer, 2020, 146, 3281-3293.	2.3	35
26	Three functional variants were identified to affect RPS24 expression and significantly associated with risk of colorectal cancer. Archives of Toxicology, 2020, 94, 295-303.	1.9	10
27	Trans-acting non-synonymous variant of FOXA1 predisposes to hepatocellular carcinoma through modulating FOXA1-ERα transcriptional program and may have undergone natural selection. Carcinogenesis, 2020, 41, 146-158.	1.3	3
28	Metabolically healthy obesity increased diabetes incidence in a middleâ€aged and elderly Chinese population. Diabetes/Metabolism Research and Reviews, 2020, 36, e3202.	1.7	21
29	LINC01149 variant modulates MICA expression that facilitates hepatitis B virus spontaneous recovery but increases hepatocellular carcinoma risk. Oncogene, 2020, 39, 1944-1956.	2.6	13
30	Evaluation of polymorphisms in microRNAâ€binding sites and pancreatic cancer risk in Chinese population. Journal of Cellular and Molecular Medicine, 2020, 24, 2252-2259.	1.6	6
31	Healthy lifestyle and cancer risk among Chinese population in the Dongfeng-Tongji cohort. Annals of Medicine, 2020, 52, 393-402.	1.5	7
32	Comparative Evaluation of Participation and Diagnostic Yield of Colonoscopy vs Fecal Immunochemical Test vs Risk-Adapted Screening in Colorectal Cancer Screening: Interim Analysis of a Multicenter Randomized Controlled Trial (TARGET-C). American Journal of Gastroenterology, 2020, 115, 1264-1274.	0.2	40
33	Challenges and recommendations for cancer care in the COVID-19 pandemic. Cancer Biology and Medicine, 2020, 17, 515-518.	1.4	7
34	Severe acute respiratory syndrome coronavirus 2 detection in the female lower genital tract. American Journal of Obstetrics and Gynecology, 2020, 223, 131-134.	0.7	58
35	Efficacy of hydrogel patches in preventing facial skin damage caused by mask compression in fighting against coronavirus disease 2019: a shortâ€term, selfâ€controlled study. Journal of the European Academy of Dermatology and Venereology, 2020, 34, e441-e443.	1.3	9
36	Functional characterization of a low-frequency V1937I variant in FASN associated with susceptibility to esophageal squamous cell carcinoma. Archives of Toxicology, 2020, 94, 2039-2046.	1.9	9

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37	Clinical Characteristics of and Medical Interventions for COVID-19 in Hemodialysis Patients in Wuhan, China. Journal of the American Society of Nephrology: JASN, 2020, 31, 1387-1397.	3.0	206
38	Challenges and countermeasures in the prevention of nosocomial infections of SARS-CoV-2 before resumption of work: Implications for the dermatology department. Journal of the American Academy of Dermatology, 2020, 83, 961-963.	0.6	1
39	Clinical characteristics and risk factors associated with COVID-19 disease severity in patients with cancer in Wuhan, China: a multicentre, retrospective, cohort study. Lancet Oncology, The, 2020, 21, 893-903.	5.1	421
40	Potential of Arbidol for Post-exposure Prophylaxis of COVID-19 Transmission: A Preliminary Report of a Retrospective Cohort Study. Current Medical Science, 2020, 40, 480-485.	0.7	39
41	Skin damage among health care workers managing coronavirus disease-2019. Journal of the American Academy of Dermatology, 2020, 82, 1215-1216.	0.6	399
42	Clinical characteristics of 54 medical staff with COVIDâ€19: A retrospective study in a single center in Wuhan, China. Journal of Medical Virology, 2020, 92, 807-813.	2.5	153
43	Risk SNP-Mediated Enhancer–Promoter Interaction Drives Colorectal Cancer through Both <i>FADS2</i> and <i>AP002754.2</i> Cancer Research, 2020, 80, 1804-1818.	0.4	50
44	Genetic variants in m6A modification genes are associated with esophageal squamous-cell carcinoma in the Chinese population. Carcinogenesis, 2020, 41, 761-768.	1.3	35
45	A functional variant in TNXB promoter associates with the risk of esophageal squamousâ€cell carcinoma. Molecular Carcinogenesis, 2020, 59, 439-446.	1.3	7
46	Neurologic Manifestations of Hospitalized Patients With Coronavirus Disease 2019 in Wuhan, China. JAMA Neurology, 2020, 77, 683.	4.5	5,308
47	N <sup>6</sup> -methyladenosine mRNA methylation of <i>PIK3CB</i> regulates AKT signalling to promote PTEN-deficient pancreatic cancer progression. Gut, 2020, 69, 2180-2192.	6.1	52
48	Serologic Detection of SARS-CoV-2 Infections in Hemodialysis Centers: A Multicenter Retrospective Study in Wuhan, China. American Journal of Kidney Diseases, 2020, 76, 490-499.e1.	2.1	33
49	Genetic Predisposition to Colon and Rectal Adenocarcinoma Is Mediated by a Super-enhancer Polymorphism Coactivating <i>CD9 </i> and <i>PLEKHG6 </i> Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 850-859.	1.1	8
50	Safety of biologics for psoriasis patients during the COVID-19 pandemic: the experience from Wuhan, China. European Journal of Dermatology, 2020, 30, 738-740.	0.3	3
51	A functional variant in the boundary of a topological association domain is associated with pancreatic cancer risk. Molecular Carcinogenesis, 2019, 58, 1855-1862.	1.3	13
52	Systematic Functional Interrogation of Genes in GWAS Loci Identified ATF1 as a Key Driver in Colorectal Cancer Modulated by a Promoter-Enhancer Interaction. American Journal of Human Genetics, 2019, 105, 29-47.	2.6	41
53	A genetic variant in <i>PIK3R1</i> is associated with pancreatic cancer survival in the Chinese population. Cancer Medicine, 2019, 8, 3575-3582.	1.3	9
54	Serum alanine transaminase levels predict type 2 diabetes risk among a middle-aged and elderly Chinese population. Annals of Hepatology, 2019, 18, 298-303.	0.6	10

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55	Association between resting heart rate and incident diabetes risk: a Mendelian randomization study. Acta Diabetologica, 2019, 56, 1037-1044.	1.2	12
56	Circulating essential metals and lung cancer: Risk assessment and potential molecular effects. Environment International, 2019, 127, 685-693.	4.8	41
57	A missense variant in PTPN12 associated with the risk of colorectal cancer by modifying Ras/MEK/ERK signaling. Cancer Epidemiology, 2019, 59, 109-114.	0.8	9
58	AWESOME: a database of SNPs that affect protein post-translational modifications. Nucleic Acids Research, 2019, 47, D874-D880.	6.5	53
59	Pancan-meQTL: a database to systematically evaluate the effects of genetic variants on methylation in human cancer. Nucleic Acids Research, 2019, 47, D1066-D1072.	6.5	45
60	Reply to comment: Serum bilirubin concentrations, type 2 diabetes, and incident coronary heart disease. Acta Diabetologica, 2019, 56, 383-384.	1.2	2
61	A functional variant rs1537373 in 9p21.3 region is associated with pancreatic cancer risk. Molecular Carcinogenesis, 2019, 58, 760-766.	1.3	9
62	Urinary levels of bisphenol A, F and S and markers of oxidative stress among healthy adult men: Variability and association analysis. Environment International, 2019, 123, 301-309.	4.8	117
63	CancerSplicingQTL: a database for genome-wide identification of splicing QTLs in human cancer. Nucleic Acids Research, 2019, 47, D909-D916.	6.5	61
64	Exome-wide analyses identify low-frequency variant in CYP26B1 and additional coding variants associated with esophageal squamous cell carcinoma. Nature Genetics, 2018, 50, 338-343.	9.4	75
65	Victims of Chinese famine in early life have increased risk of metabolic syndrome in adulthood. Nutrition, 2018, 53, 20-25.	1.1	18
66	Integrative expression quantitative trait locus–based analysis of colorectal cancer identified a functional polymorphism regulating SLC22A5 expression. European Journal of Cancer, 2018, 93, 1-9.	1.3	47
67	Association of co-exposure to heavy metals with renal function in a hypertensive population. Environment International, 2018, 112, 198-206.	4.8	41
68	Inverse association between plasma homocysteine concentrations and type 2 diabetes mellitus among a middle-aged and elderly Chinese population. Nutrition, Metabolism and Cardiovascular Diseases, 2018, 28, 278-284.	1.1	3
69	A polymorphic MYC response element in KBTBD11 influences colorectal cancer risk, especially in interaction with an MYC-regulated SNP rs6983267. Annals of Oncology, 2018, 29, 632-639.	0.6	49
70	Genetic correction of serum <scp>AFP</scp> level improves risk prediction of primary hepatocellular carcinoma in the Dongfeng–Tongji cohort study. Cancer Medicine, 2018, 7, 2691-2698.	1.3	3
71	PancanQTL: systematic identification of cis-eQTLs and trans-eQTLs in 33 cancer types. Nucleic Acids Research, 2018, 46, D971-D976.	6.5	191
72	Associations between daily cooking duration and the prevalence of diabetes and prediabetes in a middle-aged and elderly Chinese population: A cross-sectional study. Indoor Air, 2018, 28, 238-246.	2.0	10

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73	Development of a new scoring system to predict 5-year incident diabetes risk in middle-aged and older Chinese. Acta Diabetologica, 2018, 55, 13-19.	1.2	9
74	Associations of environmental exposure to metals with the risk of hypertension in China. Science of the Total Environment, 2018, 622-623, 184-191.	3.9	42
75	Environmental exposure to metals and the risk of hypertension: A cross-sectional study in China. Environmental Pollution, 2018, 233, 670-678.	3.7	70
76	Identification of Genes Associated With Hirschsprung Disease, Based on Whole-Genome Sequence Analysis, and Potential Effects on Enteric Nervous System Development. Gastroenterology, 2018, 155, 1908-1922.e5.	0.6	61
77	The advances of genetics research on Hirschsprung's disease. Pediatric Investigation, 2018, 2, 189-195.	0.6	4
78	Inhibition of kinesin family member 20B sensitizes hepatocellular carcinoma cell to microtubuleâ€targeting agents by blocking cytokinesis. Cancer Science, 2018, 109, 3450-3460.	1.7	21
79	Effect modification of CPY2E1 and GSTZ1 genetic polymorphisms on associations between prenatal disinfection by-products exposure and birth outcomes. Environmental Pollution, 2018, 243, 1126-1133.	3.7	13
80	Using different anthropometric indices to assess prediction ability of type 2 diabetes in elderly population: a 5Âyear prospective study. BMC Geriatrics, 2018, 18, 218.	1.1	38
81	Exome-wide analysis identifies three low-frequency missense variants associated with pancreatic cancer risk in Chinese populations. Nature Communications, 2018, 9, 3688.	5.8	32
82	BMI, Waist Circumference and All-Cause Mortality in a Middle-Aged and Elderly Chinese Population. Journal of Nutrition, Health and Aging, 2018, 22, 975-981.	1.5	26
83	Bidirectional association between nonalcoholic fatty liver disease and hypertension from the Dongfeng-Tongji cohort study. Journal of the American Society of Hypertension, 2018, 12, 660-670.	2.3	15
84	Educational and Behavioral Counseling in a Methadone Maintenance Treatment Program in China: A Randomized Controlled Trial. Frontiers in Psychiatry, 2018, 9, 113.	1.3	9
85	A Rare Missense Variant in TCF7L2 Associates with Colorectal Cancer Risk by Interacting with a GWAS-Identified Regulatory Variant in the MYC Enhancer. Cancer Research, 2018, 78, 5164-5172.	0.4	54
86	Prenatal phthalate exposure, birth outcomes and DNA methylation of Alu and LINE-1 repetitive elements: A pilot study in China. Chemosphere, 2018, 206, 759-765.	4.2	30
87	Reducing protein regulator of cytokinesis $1$ as a prospective therapy for hepatocellular carcinoma. Cell Death and Disease, $2018,9,534.$	2.7	48
88	A Rare Variant P507L in TPP1 Interrupts TPP1â€"TIN2 Interaction, Influences Telomere Length, and Confers Colorectal Cancer Risk in Chinese Population. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 1029-1035.	1.1	41
89	Genetic correction improves prediction efficiency of serum tumor biomarkers on digestive cancer risk in the elderly Chinese cohort study. Oncotarget, 2018, 9, 7389-7397.	0.8	7
90	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. Gut, 2017, 66, 581-587.	6.1	68

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91	Association of regular physical activity with total and cause-specific mortality among middle-aged and older Chinese: a prospective cohort study. Scientific Reports, 2017, 7, 39939.	1.6	19
92	Epigenomic landscape of 5-hydroxymethylcytosine reveals its transcriptional regulation of lncRNAs in colorectal cancer. British Journal of Cancer, 2017, 116, 658-668.	2.9	38
93	A lowâ€frequency variant in SMAD7 modulates TGFâ€Î² signaling and confers risk for colorectal cancer in Chinese population. Molecular Carcinogenesis, 2017, 56, 1798-1807.	1.3	48
94	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. Gastroenterology, 2017, 152, 2011-2021.	0.6	58
95	Serum bilirubin levels and risk of type 2 diabetes: results from two independent cohorts in middle-aged and elderly Chinese. Scientific Reports, 2017, 7, 41338.	1.6	20
96	Exposure to the Chinese famine in early life and hypertension prevalence risk in adults. Journal of Hypertension, 2017, 35, 63-68.	0.3	41
97	ELABELA and an ELABELA Fragment Protect against AKI. Journal of the American Society of Nephrology: JASN, 2017, 28, 2694-2707.	3.0	101
98	Glyceraldehydeâ€3â€phosphate dehydrogenase promotes liver tumorigenesis by modulating phosphoglycerate dehydrogenase. Hepatology, 2017, 66, 631-645.	3.6	70
99	Serum bilirubin concentrations and incident coronary heart disease risk among patients with type 2 diabetes: the Dongfeng–Tongji cohort. Acta Diabetologica, 2017, 54, 257-264.	1.2	14
100	Disparities of time trends and birth cohort effects on invasive breast cancer incidence in Shanghai and Hong Kong pre- and post-menopausal women. BMC Cancer, 2017, 17, 362.	1.1	5
101	A functional variant in GREM1 confers risk for colorectal cancer by disrupting a hsa-miR-185-3p binding site. Oncotarget, 2017, 8, 61318-61326.	0.8	20
102	Bidirectional association between nonalcoholic fatty liver disease and type 2 diabetes in Chinese population: Evidence from the Dongfeng-Tongji cohort study. PLoS ONE, 2017, 12, e0174291.	1.1	48
103	Independent and joint effects of moderate alcohol consumption and smoking on the risks of non-alcoholic fatty liver disease in elderly Chinese men. PLoS ONE, 2017, 12, e0181497.	1.1	28
104	BRCA1 missense polymorphisms are associated with poor prognosis of pancreatic cancer patients in a Chinese population. Oncotarget, 2017, 8, 36033-36039.	0.8	21
105	The Relationship between Serum Bilirubin and Elevated Fibrotic Indices among HBV Carriers: A Cross-Sectional Study of a Chinese Population. International Journal of Molecular Sciences, 2016, 17, 2057.	1.8	12
106	A functional polymorphism located at transcription factor binding sites, rs6695837 near LAMC1 gene, confers risk of colorectal cancer in Chinese populations. Carcinogenesis, 2016, 38, bgw204.	1.3	59
107	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. Nature Communications, 2016, 7, 11478.	5.8	73
108	Sleep Duration and Midday Napping with 5-Year Incidence and Reversion of Metabolic Syndrome in Middle-Aged and Older Chinese. Sleep, 2016, 39, 1911-1918.	0.6	35

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109	A novel variant on chromosome 6p21.1 is associated with the risk of developing colorectal cancer: a two-stage case-control study in Han Chinese. BMC Cancer, 2016, 16, 807.	1.1	1
110	Association between serum bilirubin levels and decline in estimated glomerular filtration rate among patients with type 2 diabetes. Journal of Diabetes and Its Complications, 2016, 30, 1255-1260.	1.2	16
111	Pancreatic cancer risk variant in LINC00673 creates a miR-1231 binding site and interferes with PTPN11 degradation. Nature Genetics, 2016, 48, 747-757.	9.4	237
112	Association of shift-work, daytime napping, and nighttime sleep with cancer incidence and cancer-caused mortality in Dongfeng-tongji cohort study. Annals of Medicine, 2016, 48, 641-651.	1.5	22
113	Exposure to the Chinese Famine in Childhood Increases Type 2 Diabetes Risk in Adults. Journal of Nutrition, 2016, 146, 2289-2295.	1.3	70
114	Green tea consumption is associated with reduced incident CHD and improved CHD-related biomarkers in the Dongfeng-Tongji cohort. Scientific Reports, 2016, 6, 24353.	1.6	34
115	Association between bilirubin and risk of Non-Alcoholic Fatty Liver Disease based on a prospective cohort study. Scientific Reports, 2016, 6, 31006.	1.6	39
116	Longer Sleep Duration and Midday Napping Are Associated with a Higher Risk of CHD Incidence in Middle-Aged and Older Chinese: the Dongfeng-Tongji Cohort Study. Sleep, 2016, 39, 645-652.	0.6	64
117	Nighttime sleep duration and risk of nonalcoholic fatty liver disease: the Dongfeng-Tongji prospective study. Annals of Medicine, 2016, 48, 468-476.	1.5	19
118	<scp><i>Helicobacter pylori</i></scp> infection is associated with type 2 diabetes among a middleâ€and oldâ€age Chinese population. Diabetes/Metabolism Research and Reviews, 2016, 32, 95-101.	1.7	43
119	Long sleep duration and afternoon napping are associated with higher risk of incident diabetes in middle-aged and older Chinese: the Dongfeng-Tongji cohort study. Annals of Medicine, 2016, 48, 216-223.	1.5	34
120	A functional polymorphism in <i>lnc-LAMC2-1:1</i> confers risk of colorectal cancer by affecting miRNA binding. Carcinogenesis, 2016, 37, 443-451.	1.3	68
121	A single nucleotide polymorphism in the $3\hat{a}\in^2$ -UTR of STAT3 regulates its expression and reduces risk of pancreatic cancer in a Chinese population. Oncotarget, 2016, 7, 62305-62311.	0.8	10
122	A functional variant rs4442975 modulating FOXA1-binding affinity does not influence the risk or progression of breast cancer in Chinese Han population. Oncotarget, 2016, 7, 81691-81697.	0.8	3
123	Identification of a functional variant for colorectal cancer risk mapping to chromosome 5q31.1. Oncotarget, 2016, 7, 35199-35207.	0.8	12
124	MAD1L1 Arg558His and MAD2L1 Leu84Met interaction with smoking increase the risk of colorectal cancer. Scientific Reports, 2015, 5, 12202.	1.6	20
125	Systematic Confirmation Study of GWAS-Identified Genetic Variants for Kawasaki Disease in A Chinese Population. Scientific Reports, 2015, 5, 8194.	1.6	31
126	Dietary legume consumption reduces risk of colorectal cancer: evidence from a meta-analysis of cohort studies. Scientific Reports, 2015, 5, 8797.	1.6	79

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127	Association of Adiposity Indices with Platelet Distribution Width and Mean Platelet Volume in Chinese Adults. PLoS ONE, 2015, 10, e0129677.	1.1	9
128	Association between polymorphisms in cdc27 and breast cancer in a Chinese population. Tumor Biology, 2015, 36, 5299-5304.	0.8	5
129	A Solute Carrier Family 22 Member 3 Variant rs3088442 G→A Associated with Coronary Heart Disease Inhibits Lipopolysaccharide-induced Inflammatory Response. Journal of Biological Chemistry, 2015, 290, 5328-5340.	1.6	34
130	Housing Characteristics in Relation to Exhaled Nitric Oxide in China. American Journal of Health Behavior, 2015, 39, 88-98.	0.6	6
131	CYP2C19 polymorphism and clinical outcomes among patients of different races treated with clopidogrel: A systematic review and meta-analysis. Journal of Huazhong University of Science and Technology [Medical Sciences], 2015, 35, 147-156.	1.0	37
132	IncRNASNP: a database of SNPs in IncRNAs and their potential functions in human and mouse. Nucleic Acids Research, 2015, 43, D181-D186.	6.5	204
133	Two Novel Variants on 13q22.1 Are Associated with Risk of Esophageal Squamous Cell Carcinoma. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1774-1780.	1.1	12
134	Serum creatinine levels and risk of metabolic syndrome in a middle-aged and older Chinese population. Clinica Chimica Acta, 2015, 440, 177-182.	0.5	13
135	Meta-analysis of genome-wide association studies of adult height in East Asians identifies 17 novel loci. Human Molecular Genetics, 2015, 24, 1791-1800.	1.4	105
136	Genetic variants in the SWI/SNF complex and smoking collaborate to modify the risk of pancreatic cancer in a Chinese population. Molecular Carcinogenesis, 2015, 54, 761-768.	1.3	35
137	A Phosphorylation-Related Variant ADD1-rs4963 Modifies the Risk of Colorectal Cancer. PLoS ONE, 2015, 10, e0121485.	1.1	8
138	SF3A1 and pancreatic cancer: new evidence for the association of the spliceosome and cancer. Oncotarget, 2015, 6, 37750-37757.	0.8	20
139	New Integrated Strategy Emphasizing Infection Source Control to Curb Schistosomiasis japonica in a Marshland Area of Hubei Province, China: Findings from an Eight-Year Longitudinal Survey. PLoS ONE, 2014, 9, e89779.	1.1	23
140	Dietary Mushroom Intake May Reduce the Risk of Breast Cancer: Evidence from a Meta-Analysis of Observational Studies. PLoS ONE, 2014, 9, e93437.	1.1	40
141	A Genetic Variant rs1801274 in FCGR2A as a Potential Risk Marker for Kawasaki Disease: A Case-Control Study and Meta-Analysis. PLoS ONE, 2014, 9, e103329.	1.1	32
142	Integrative Genomic Analysis Identifies That SERPINA6-rs1998056 Regulated by FOXA/ERα Is Associated with Female Hepatocellular Carcinoma. PLoS ONE, 2014, 9, e107246.	1.1	9
143	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. Human Molecular Genetics, 2014, 23, 6616-6633.	1.4	90
144	A genome wide association study of genetic loci that influence tumour biomarkers cancer antigen 19-9, carcinoembryonic antigen and $l_{\pm}$ fetoprotein and their associations with cancer risk. Gut, 2014, 63, 143-151.	6.1	67

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145	Association between polymorphisms within the susceptibility region 8q24 and breast cancer in a Chinese population. Tumor Biology, 2014, 35, 2649-2654.	0.8	9
146	Reelin gene variants and risk of autism spectrum disorders: An integrated metaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 192-200.	1.1	85
147	Variants in the 5′-upstream region of GPC5 confer risk of lung cancer in never smokers. Cancer Epidemiology, 2014, 38, 66-72.	0.8	11
148	A genetic variant in microRNA target site of TGF- $\hat{l}^2$ signaling pathway increases the risk of colorectal cancer in a Chinese population. Tumor Biology, 2014, 35, 4301-4306.	0.8	14
149	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. Nature Genetics, 2014, 46, 1001-1006.	9.4	148
150	Allium Vegetables and Garlic Supplements Do Not Reduce Risk of Colorectal Cancer, Based on Meta-analysis of Prospective Studies. Clinical Gastroenterology and Hepatology, 2014, 12, 1991-2001.e4.	2.4	53
151	Non-linear dose–response relationship between cigarette smoking and pancreatic cancer risk: Evidence from a meta-analysis of 42 observational studies. European Journal of Cancer, 2014, 50, 193-203.	1.3	63
152	Metaâ€analysis on night shift work and risk of metabolic syndrome. Obesity Reviews, 2014, 15, 709-720.	3.1	218
153	A Functional p53 Responsive Polymorphism in KITLG, rs4590952, Does not Affect the Risk of Breast Cancer. Scientific Reports, 2014, 4, 6371.	1.6	4
154	Parity and pancreatic cancer risk: evidence from a meta-analysis of twenty epidemiologic studies. Scientific Reports, 2014, 4, 5313.	1.6	20
155	8p22–23-rs2254546 as a Susceptibility Locus for Kawasaki Disease: a Case-control Study and a Meta-analysis. Scientific Reports, 2014, 4, 4247.	1.6	11
156	The Roles of Ca2+/NFAT Signaling Genes in Kawasaki Disease: Single- and Multiple-Risk Genetic Variants. Scientific Reports, 2014, 4, 5208.	1.6	15
157	Genetic variant in SWI/SNF complexes influences hepatocellular carcinoma risk: a new clue for the contribution of chromatin remodeling in carcinogenesis. Scientific Reports, 2014, 4, 4147.	1.6	23
158	Multiâ€loci analysis reveals the importance of genetic variations in sensitivity of platinumâ€based chemotherapy in nonâ€smallâ€cell lung cancer. Molecular Carcinogenesis, 2013, 52, 923-931.	1.3	25
159	Association between serum uric acid and the metabolic syndrome among a middle- and old-age Chinese population. European Journal of Epidemiology, 2013, 28, 669-676.	2.5	72
160	Exome Sequencing Identified NRG3 as a Novel Susceptible Gene of Hirschsprung's Disease in a Chinese Population. Molecular Neurobiology, 2013, 47, 957-966.	1.9	30
161	Risk of GWAS-identified genetic variants for breast cancer in a Chinese population: a multiple interaction analysis. Breast Cancer Research and Treatment, 2013, 142, 637-644.	1.1	18
162	Common genetic variants regulating ADD3 gene expression alter biliary atresia risk. Journal of Hepatology, 2013, 59, 1285-1291.	1.8	84

#	Article	IF	CITATIONS
163	Genome-wide copy number variation study in anorectal malformations. Human Molecular Genetics, 2013, 22, 621-631.	1.4	21
164	Meta-analysis of the Association Between DCDC2 Polymorphisms and Risk of Dyslexia. Molecular Neurobiology, 2013, 47, 435-442.	1.9	24
165	Identification of common variants in BRCA2 and MAP2K4 for susceptibility to sporadic pancreatic cancer. Carcinogenesis, 2013, 34, 1001-1005.	1.3	19
166	GSK-3β Polymorphism Discriminates Bipolar Disorder and Schizophrenia: A Systematic Meta-Analysis. Molecular Neurobiology, 2013, 48, 404-411.	1.9	29
167	Oncogenic B-RafV600E abrogates the AKT/B-Raf/Mps1 interaction in melanoma cells. Cancer Letters, 2013, 337, 125-132.	3.2	15
168	A genome-wide association study identifies two new cervical cancer susceptibility loci at 4q12 and 17q12. Nature Genetics, 2013, 45, 918-922.	9.4	108
169	Association of IL-1 Receptor Antagonist Gene VNTR Polymorphism with Ischemic Stroke in the Chinese Uyghur Population. Biochemical Genetics, 2013, 51, 698-706.	0.8	7
170	Cohort Profile: The Dongfeng–Tongji cohort study of retired workers. International Journal of Epidemiology, 2013, 42, 731-740.	0.9	219
171	Risk prediction of esophageal squamous-cell carcinoma with common genetic variants and lifestyle factors in Chinese population. Carcinogenesis, 2013, 34, 1782-1786.	1.3	37
172	Genetic variations in the TGFÂ signaling pathway, smoking and risk of colorectal cancer in a Chinese population. Carcinogenesis, 2013, 34, 936-942.	1.3	69
173	The Caseâ€Only Test for Gene–Environment Interaction is Not Uniformly Powerful: An Empirical Example. Genetic Epidemiology, 2013, 37, 402-407.	0.6	8
174	Allele-Specific Expression at the <i>RET </i> Locus in Blood and Gut Tissue of Individuals Carrying Risk Alleles for Hirschsprung Disease. Human Mutation, 2013, 34, 754-762.	1.1	4
175	Genetic variations in <i>TERTâ€CLPTM1111</i>	1.3	44
176	Different Physical Activity Subtypes and Risk of Metabolic Syndrome in Middle-Aged and Older Chinese People. PLoS ONE, 2013, 8, e53258.	1.1	36
177	Prevalence and Associated Risk Factors of Dyslexic Children in a Middle-Sized City of China: A Cross-Sectional Study. PLoS ONE, 2013, 8, e56688.	1.1	55
178	Genetic Variants in the Folate Pathway and the Risk of Neural Tube Defects: A Meta-Analysis of the Published Literature. PLoS ONE, 2013, 8, e59570.	1.1	74
179	The Leptin Gene Family and Colorectal Cancer: Interaction with Smoking Behavior and Family History of Cancer. PLoS ONE, 2013, 8, e60777.	1.1	23
180	Replication Study in Chinese Population and Meta-Analysis Supports Association of the 5p15.33 Locus with Lung Cancer. PLoS ONE, 2013, 8, e62485.	1.1	17

#	Article	IF	Citations
181	Associations between Two Genetic Variants in NKX2-5 and Risk of Congenital Heart Disease in Chinese Population: A Meta-Analysis. PLoS ONE, 2013, 8, e70979.	1.1	14
182	Functional Polymorphisms in FAS/FASL System Increase the Risk of Neuroblastoma in Chinese Population. PLoS ONE, 2013, 8, e71656.	1.1	30
183	Application of a Hybrid Model for Predicting the Incidence of Tuberculosis in Hubei, China. PLoS ONE, 2013, 8, e80969.	1.1	27
184	A Genome Wide Association Study Identifies Common Variants Associated with Lipid Levels in the Chinese Population. PLoS ONE, 2013, 8, e82420.	1.1	57
185	The Genetic Variant on Chromosome 10p14 Is Associated with Risk of Colorectal Cancer: Results from a Case-Control Study and a Meta-Analysis. PLoS ONE, 2013, 8, e64310.	1.1	9
186	MNS16A Tandem Repeats Minisatellite of Human Telomerase Gene and Cancer Risk: A Meta-Analysis. PLoS ONE, 2013, 8, e73367.	1.1	8
187	The SNP rs402710 in 5p15.33 Is Associated with Lung Cancer Risk: A Replication Study in Chinese Population and a Meta-Analysis. PLoS ONE, 2013, 8, e76252.	1.1	13
188	A Functional Variant rs1820453 in YAP1 and Breast Cancer Risk in Chinese Population. PLoS ONE, 2013, 8, e79056.	1.1	9
189	HBV-Related Hepatocellular Carcinoma Susceptibility Gene KIF1B Is Not Associated with Development of Chronic Hepatitis B. PLoS ONE, 2012, 7, e28839.	1.1	21
190	Genome-Wide Copy Number Analysis Uncovers a New HSCR Gene: NRG3. PLoS Genetics, 2012, 8, e1002687.	1.5	51
191	Genetic Variants at 6p21.1 and 7p15.3 Are Associated with Risk of Multiple Cancers in Han Chinese. American Journal of Human Genetics, 2012, 91, 928-934.	2.6	76
192	Genetic variant in <i>KIAA0319</i> , but not in <i>DYX1C1</i> , is associated with risk of dyslexia: An integrated metaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 970-976.	1.1	38
193	The SLC4A7 variant rs4973768 is associated with breast cancer risk: evidence from a case–control study and a meta-analysis. Breast Cancer Research and Treatment, 2012, 136, 847-857.	1.1	39
194	Genome-wide association analyses of esophageal squamous cell carcinoma in Chinese identify multiple susceptibility loci and gene-environment interactions. Nature Genetics, 2012, 44, 1090-1097.	9.4	238
195	Genome-wide association study identifies five loci associated with susceptibility to pancreatic cancer in Chinese populations. Nature Genetics, 2012, 44, 62-66.	9.4	164
196	Polymorphisms in TP53 and MDM2 contribute to higher risk of colorectal cancer in Chinese population: a hospital-based, case–control study. Molecular Biology Reports, 2012, 39, 9661-9668.	1.0	15
197	A Common SMAD7 Variant Is Associated with Risk of Colorectal Cancer: Evidence from a Case-Control Study and a Meta-Analysis. PLoS ONE, 2012, 7, e33318.	1.1	21
198	The SNP rs961253 in 20p12.3 Is Associated with Colorectal Cancer Risk: A Case-Control Study and a Meta-Analysis of the Published Literature. PLoS ONE, 2012, 7, e34625.	1.1	14

#	Article	IF	CITATIONS
199	The epidemiology and etiology of influenzaâ€like illness in Chinese children from 2008 to 2010. Journal of Medical Virology, 2012, 84, 672-678.	2.5	26
200	Mutations in the NRG1 gene are associated with Hirschsprung disease. Human Genetics, 2012, 131, 67-76.	1.8	51
201	The Expression Levels of Plasma micoRNAs in Atrial Fibrillation Patients. PLoS ONE, 2012, 7, e44906.	1.1	97
202	Association of candidate genetic variations with gastric cardia adenocarcinoma in Chinese population: a multiple interaction analysis. Carcinogenesis, 2011, 32, 336-342.	1.3	45
203	Genome-wide association study identifies three new susceptibility loci for esophageal squamous-cell carcinoma in Chinese populations. Nature Genetics, 2011, 43, 679-684.	9.4	260
204	Assessment of XPD Lys751Gln and XRCC1 T–77C polymorphisms in advanced non-small-cell lung cancer patients treated with platinum-based chemotherapy. Lung Cancer, 2011, 73, 110-115.	0.9	31
205	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. Nature Genetics, 2011, 43, 1215-1218.	9.4	250
206	The effect of laparoscopic excision vs open excision in children with choledochal cyst: a midterm follow-up study. Journal of Pediatric Surgery, 2011, 46, 662-665.	0.8	61
207	Interactions between Genetic Variants in the Adiponectin, Adiponectin Receptor 1 and Environmental Factors on the Risk of Colorectal Cancer. PLoS ONE, 2011, 6, e27301.	1.1	41
208	RET Mutational Spectrum in Hirschsprung Disease: Evaluation of 601 Chinese Patients. PLoS ONE, 2011, 6, e28986.	1.1	26
209	No NRG1 V266L in Chinese patients with schizophrenia. Psychiatric Genetics, 2011, 21, 47-49.	0.6	2
210	Combined Effect of Genetic Polymorphisms in P53, P73, and MDM2 on Non-small Cell Lung Cancer Survival. Journal of Thoracic Oncology, 2011, 6, 1793-1800.	0.5	38
211	A functional â^777T>C polymorphism in XRCC1 is associated with risk of breast cancer. Breast Cancer Research and Treatment, 2011, 125, 479-487.	1.1	32
212	Interaction of <i>Cyclooxygenaseâ€2</i> promoter polymorphisms with <i>Helicobacter pylori</i> infection and risk of gastric cancer. Molecular Carcinogenesis, 2011, 50, 876-883.	1.3	18
213	Association between polymorphisms of <i>DRD2</i> and <i>DRD4</i> and opioid dependence: Evidence from the current studies. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 661-670.	1.1	55
214	A functional polymorphism (â^'1607 1Gâ†'2G) in the matrix metalloproteinase†promoter is associated with development and progression of lung cancer. Cancer, 2011, 117, 5172-5181.	2.0	35
215	Variants of <i>HSPA1A </i> in Combination with Plasma Hsp70 and Anti-Hsp70 Antibody Levels Associated with Higher Risk of Acute Coronary Syndrome. Cardiology, 2011, 119, 57-64.	0.6	9
216	Variations in <i>HSPA1B</i> at 6p21.3 Are Associated with Lung Cancer Risk and Prognosis in Chinese Populations. Cancer Research, 2011, 71, 7576-7586.	0.4	30

#	Article	IF	Citations
217	Fine Mapping of the NRG1 Hirschsprung's Disease Locus. PLoS ONE, 2011, 6, e16181.	1.1	37
218	Evaluation of Genetic Susceptibility Loci for Chronic Hepatitis B in Chinese: Two Independent Case-Control Studies. PLoS ONE, 2011, 6, e17608.	1.1	69
219	Smoking and COX-2 Functional Polymorphisms Interact to Increase the Risk of Gastric Cardia Adenocarcinoma in Chinese Population. PLoS ONE, 2011, 6, e21894.	1.1	22
220	An Integrated Meta-Analysis of Two Variants in HOXA1/HOXB1 and Their Effect on the Risk of Autism Spectrum Disorders. PLoS ONE, 2011, 6, e25603.	1.1	9
221	Differential Contributions of Rare and Common, Coding and Noncoding Ret Mutations to Multifactorial Hirschsprung Disease Liability. American Journal of Human Genetics, 2010, 87, 60-74.	2.6	230
222	Fine mapping of the 9q31 Hirschsprung's disease locus. Human Genetics, 2010, 127, 675-683.	1.8	27
223	Polymorphisms of methylenetetrahydrofolate reductase are associated with a high risk of nasopharyngeal carcinoma in a smoking population from southern China. Molecular Carcinogenesis, 2010, 49, 928-934.	1.3	16
224	Polymorphisms of death pathway genes FAS and FASL and risk of nasopharyngeal carcinoma. Molecular Carcinogenesis, 2010, 49, 944-950.	1.3	27
225	Haplotype Analysis Reveals a Possible Founder Effect of RET Mutation R114H for Hirschsprung's Disease in the Chinese Population. PLoS ONE, 2010, 5, e10918.	1.1	19
226	Reduced RET expression in gut tissue of individuals carrying risk alleles of Hirschsprung's disease. Human Molecular Genetics, 2010, 19, 1461-1467.	1.4	47
227	Genome-wide association study identifies a susceptibility locus for biliary atresia on 10q24.2. Human Molecular Genetics, 2010, 19, 2917-2925.	1.4	117
228	Functional evaluation of missense variations in the human MAD1L1 and MAD2L1 genes and their impact on susceptibility to lung cancer. Journal of Medical Genetics, 2010, 47, 616-622.	1.5	42
229	Genome-Wide Interrogation Identifies <i>YAP1</i> Variants Associated with Survival of Small-Cell Lung Cancer Patients. Cancer Research, 2010, 70, 9721-9729.	0.4	53
230	ICPS: an integrative cancer profiler system. Bioinformatics, 2010, 26, 2649-2650.	1.8	2
231	No association betweenXRCC1polymorphisms and survival in non-small-cell lung cancer patients treated with platinum-based chemotherapy. Cancer Biology and Therapy, 2010, 10, 854-859.	1.5	12
232	Lack of association between nNOS $\hat{a}^3$ 84G>A polymorphism and risk of infantile hypertrophic pyloric stenosis in a Chinese population. Journal of Pediatric Surgery, 2010, 45, 709-713.	0.8	10
233	Genome-wide association study identifies <i>NRG1</i> as a susceptibility locus for Hirschsprung's disease. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 2694-2699.	3.3	171
234	MNX1 (HLXB9) mutations in Currarino patients. Journal of Pediatric Surgery, 2009, 44, 1892-1898.	0.8	33

#	Article	IF	CITATIONS
235	Mutational analysis of <i>SHH</i> and <i>GLI3</i> in anorectal malformations. Birth Defects Research Part A: Clinical and Molecular Teratology, 2008, 82, 644-648.	1.6	17
236	Identification of a <i>HOXD13</i> mutation in a VACTERL patient. American Journal of Medical Genetics, Part A, 2008, 146A, 3181-3185.	0.7	82
237	Mapping of a Hirschsprung's disease locus in 3p21. European Journal of Human Genetics, 2008, 16, 833-840.	1.4	16
238	Perturbation of Hoxb5 Signaling in Vagal Neural Crests Down-Regulates Ret Leading to Intestinal Hypoganglionosis in Mice. Gastroenterology, 2008, 134, 1104-1115.	0.6	49
239	Functional Variants in Cell Death Pathway Genes and Risk of Pancreatic Cancer. Clinical Cancer Research, 2008, 14, 3230-3236.	3.2	82
240	Interaction of P53 Arg72Pro and MDM2 T309G polymorphisms and their associations with risk of gastric cardia cancer. Carcinogenesis, 2007, 28, 1996-2001.	1.3	68
241	Role of RET and ko=PHOX2B gene polymorphisms in risk of Hirschsprung's disease in Chinese population. Gut, 2007, 56, 736-736.	6.1	22
242	Implications of Endocrine Gland–Derived Vascular Endothelial Growth Factor/Prokineticin-1 Signaling in Human Neuroblastoma Progression. Clinical Cancer Research, 2007, 13, 868-875.	3.2	47
243	Hirschsprung disease, associated syndromes and genetics: a review. Journal of Medical Genetics, 2007, 45, 1-14.	1.5	848
244	Involvement of visinin-like protein-1 (VSNL-1) in regulating proliferative and invasive properties of neuroblastoma. Carcinogenesis, 2007, 28, 2122-2130.	1.3	17
245	Genetic Polymorphisms in Folate- Metabolizing Enzymes and Risk of Gastroesophageal Cancers: A Potential Nutrient-Gene Interaction in Cancer Development. Forum of Nutrition, 2007, 60, 140-145.	3.7	16
246	Platelet 12-lipoxygenase Arg261Gln polymorphism: functional characterization and association with risk of esophageal squamous cell carcinoma in combination with COX-2 polymorphisms. Pharmacogenetics and Genomics, 2007, 17, 197-205.	0.7	35
247	Application of HapMap data to the evaluation of 8 candidate genes for pediatric slow transit constipation. Journal of Pediatric Surgery, 2007, 42, 666-671.	0.8	6
248	Correlation Between Genetic Variations in Hox Clusters and Hirschsprung's Disease. Annals of Human Genetics, 2007, 71, 526-536.	0.3	27
249	Evaluation of the Thyroid Transcription Factor-1 Gene (TITF1) as a Hirschsprung's Disease Locus. Annals of Human Genetics, 2007, 71, 746-754.	0.3	13
250	Adenosine Diphosphate Ribosyl Transferase and X-Ray Repair Cross-Complementing 1 Polymorphisms in Gastric Cardia Cancer. Gastroenterology, 2006, 131, 420-427.	0.6	53
251	Integrated genome-wide gene expression map and high-resolution analysis of aberrant chromosomal regions in squamous cell lung cancer. FEBS Letters, 2006, 580, 2774-2778.	1.3	4
252	A novel T-77C polymorphism in DNA repair gene XRCC1 contributes to diminished promoter activity and increased risk of non-small cell lung cancer. Oncogene, 2006, 25, 3613-3620.	2.6	115

#	Article	IF	Citations
253	Polymorphisms of XRCC1 genes and risk of nasopharyngeal carcinoma in the Cantonese population. BMC Cancer, 2006, 6, 167.	1.1	54
254	Genetic polymorphisms in cell cycle regulatory genesMDM2 and TP53 are associated with susceptibility to lung cancer. Human Mutation, 2006, 27, 110-117.	1.1	142
255	Identification of a novel polymorphism Arg290Gln of esophageal cancer related gene 1 (ECRG1) and its related risk to esophageal squamous cell carcinoma. Carcinogenesis, 2006, 27, 798-802.	1.3	24
256	Liver Intestine-Cadherin (CDH17) Haplotype Is Associated with Increased Risk of Hepatocellular Carcinoma. Clinical Cancer Research, 2006, 12, 5248-5252.	3.2	34
257	Common genetic polymorphisms in the 5′-flanking Region of the SULT1A1 gene: haplotypes and their association with platelet enzymatic activity. Pharmacogenetics and Genomics, 2005, 15, 465-473.	0.7	37
258	Dinucleotide polymorphism of p73 gene is associated with a reduced risk of lung cancer in a Chinese population. International Journal of Cancer, 2005, 114, 455-460.	2.3	47
259	DNA repair gene XPC genotypes/haplotypes and risk of lung cancer in a Chinese population. International Journal of Cancer, 2005, 115, 478-483.	2.3	87
260	Genetic polymorphisms of interleukin (IL)-1B, IL-1RN, IL-8, IL-10 and tumor necrosis factor $\hat{l}_{\pm}$ and risk of gastric cancer in a Chinese population. Carcinogenesis, 2005, 26, 631-636.	1.3	212
261	Significant increase in risk of gastroesophageal cancer is associated with interaction between promoter polymorphisms in thymidylate synthase and serum folate status. Carcinogenesis, 2005, 26, 1430-1435.	1.3	37
262	FASL â€"844C polymorphism is associated with increased activation-induced T cell death and risk of cervical cancer. Journal of Experimental Medicine, 2005, 202, 967-974.	4.2	104
263	The Role of P53 and MDM2 Polymorphisms in the Risk of Esophageal Squamous Cell Carcinoma. Cancer Research, 2005, 65, 9582-9587.	0.4	169
264	Functional polymorphisms in cell death pathway genes FAS and FASL contribute to risk of lung cancer. Journal of Medical Genetics, 2005, 42, 479-484.	1.5	120
265	Aurora- A/STK15 T $\pm$ 91A is a general low penetrance cancer susceptibility gene: a meta-analysis of multiple cancer types. Carcinogenesis, 2005, 26, 1368-1373.	1.3	132
266	Polymorphisms and mutations of the folate receptor-alpha gene and risk of gastric cancer in a Chinese population. International Journal of Molecular Medicine, 2005, 15, 627.	1.8	0
267	Identification of Functional Genetic Variants in and Their Association With Risk of Esophageal Cancer. Gastroenterology, 2005, 129, 565-576.	0.6	100
268	Functional haplotypes in the promoter of matrix metalloproteinase-2 and lung cancer susceptibility. Carcinogenesis, 2005, 26, 1117-1121.	1.3	78
269	Identification of Functional Genetic Variants in Cyclooxygenase-2 and Their Association With Risk of Esophageal Cancer. Gastroenterology, 2005, 129, 565-576.	0.6	200
270	A common polymorphism in the 3′UTR of cyclooxygenase 2/prostaglandin synthase 2 gene and risk of lung cancer in a Chinese population. Lung Cancer, 2005, 48, 11-17.	0.9	68

#	Article	IF	CITATIONS
271	Genetic Polymorphisms in Methylenetetrahydrofolate Reductase and Thymidylate Synthase and Risk of Pancreatic Cancer. Clinical Gastroenterology and Hepatology, 2005, 3, 743-751.	2.4	56
272	Polymorphisms in DNA base excision repair genes ADPRT and XRCC1 and risk of lung cancer. Cancer Research, 2005, 65, 722-6.	0.4	127
273	Functional STK15 Phe31lle Polymorphism Is Associated with the Occurrence and Advanced Disease Status of Esophageal Squamous Cell Carcinoma. Cancer Research, 2004, 64, 2680-2683.	0.4	80
274	Functional Phe31lle polymorphism in Aurora A and risk of breast carcinoma. Carcinogenesis, 2004, 25, 2225-2230.	1.3	57
275	Functional Haplotypes in the Promoter of Matrix Metalloproteinase-2 Predict Risk of the Occurrence and Metastasis of Esophageal Cancer. Cancer Research, 2004, 64, 7622-7628.	0.4	182
276	Polymorphisms of Death Pathway Genes FAS and FASL in Esophageal Squamous-Cell Carcinoma. Journal of the National Cancer Institute, 2004, 96, 1030-1036.	3.0	185
277	Identification of Genetic Variants in Base Excision Repair Pathway and Their Associations with Risk of Esophageal Squamous Cell Carcinoma. Cancer Research, 2004, 64, 4378-4384.	0.4	208
278	Loss of myeloid-related proteins 8 and myeloid-related proteins 14 expression in human esophageal squamous cell carcinoma correlates with poor differentiation. World Journal of Gastroenterology, 2004, 10, 1093.	1.4	40
279	Sequence variations in the DNA repair geneXPD and risk of lung cancer in a Chinese population. International Journal of Cancer, 2003, 105, 669-673.	2.3	102
280	Substantial reduction in risk of breast cancer associated with genetic polymorphisms in the promoters of the matrix metalloproteinase-2 and tissue inhibitor of metalloproteinase-2 genes. Carcinogenesis, 2003, 25, 399-404.	1.3	107
281	A functional polymorphism in the SULT1A1 gene (G638A) is associated with risk of lung cancer in relation to tobacco smoking. Carcinogenesis, 2003, 25, 773-778.	1.3	51
282	Inactivation of DNA repair gene O6-methylguanine-DNA methyltransferase by promoter hypermethylation and its relation to p53 mutations in esophageal squamous cell carcinoma. Carcinogenesis, 2003, 24, 1039-1044.	1.3	53
283	A functional polymorphism in the matrix metalloproteinase-2 gene promoter (-1306C/T) is associated with risk of development but not metastasis of gastric cardia adenocarcinoma. Cancer Research, 2003, 63, 3987-90.	0.4	73
284	Polymorphisms of DNA repair genesXRCC1 and XPD and their associations with risk of esophageal squamous cell carcinoma in a Chinese population. International Journal of Cancer, 2002, 100, 600-605.	2.3	120
285	Genetic polymorphism in myeloperoxidase but not GSTM1 is associated with risk of lung squamous cell carcinoma in a Chinese population. International Journal of Cancer, 2002, 102, 275-279.	2.3	47
286	Susceptibility to gastric cardia adenocarcinoma and genetic polymorphisms in methylenetetrahydrofolate reductase in an at-risk Chinese population. Cancer Epidemiology Biomarkers and Prevention, 2002, 11, 1454-8.	1.1	34