

Qiuyin Cai

List of Publications by Year in descending order

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Version: 2024-02-01

388
papers

23,631
citations

9234

74
h-index

13727

129
g-index

397
all docs

397
docs citations

397
times ranked

30937
citing authors

#	ARTICLE	IF	CITATIONS
1	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
3	Genome-wide trans-ancestry meta-analysis provides insight into the genetic architecture of type 2 diabetes susceptibility. <i>Nature Genetics</i> , 2014, 46, 234-244.	9.4	959
4	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	9.4	513
5	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
6	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	9.4	492
7	Genome-wide association study identifies a new breast cancer susceptibility locus at 6q25.1. <i>Nature Genetics</i> , 2009, 41, 324-328.	9.4	481
8	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
9	Meta-analysis identifies common variants associated with body mass index in east Asians. <i>Nature Genetics</i> , 2012, 44, 307-311.	9.4	372
10	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	13.7	353
11	The trans-ancestral genomic architecture of glycemic traits. <i>Nature Genetics</i> , 2021, 53, 840-860.	9.4	341
12	The Human Tumor Atlas Network: Charting Tumor Transitions across Space and Time at Single-Cell Resolution. <i>Cell</i> , 2020, 181, 236-249.	13.5	334
13	The landscape of recombination in African Americans. <i>Nature</i> , 2011, 476, 170-175.	13.7	319
14	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
15	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
16	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. <i>Nature Genetics</i> , 2014, 46, 533-542.	9.4	212
17	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	2.6	201
18	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , 2011, 43, 570-573.	9.4	198

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19	Identification of New Genetic Risk Variants for Type 2 Diabetes. <i>PLoS Genetics</i> , 2010, 6, e1001127.	1.5	193
20	Meta-analysis of genome-wide association studies in East Asian-ancestry populations identifies four new loci for body mass index. <i>Human Molecular Genetics</i> , 2014, 23, 5492-5504.	1.4	192
21	Meta-Analysis of Genome-Wide Association Studies in African Americans Provides Insights into the Genetic Architecture of Type 2 Diabetes. <i>PLoS Genetics</i> , 2014, 10, e1004517.	1.5	191
22	Dietary flavonoids, quercetin, luteolin and genistein, reduce oxidative DNA damage and lipid peroxidation and quench free radicals. <i>Cancer Letters</i> , 1997, 119, 99-107.	3.2	186
23	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	9.4	184
24	Circulating 25-Hydroxyvitamin D and Risk of Pancreatic Cancer: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. <i>American Journal of Epidemiology</i> , 2010, 172, 81-93.	1.6	181
25	Urinary isothiocyanate levels, brassica, and human breast cancer. <i>Cancer Research</i> , 2003, 63, 3980-6.	0.4	175
26	Genome-wide association analyses in east Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2013, 45, 191-196.	9.4	173
27	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
28	Southern community cohort study: establishing a cohort to investigate health disparities. <i>Journal of the National Medical Association</i> , 2005, 97, 972-9.	0.6	160
29	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	7.7	157
30	Inhibition of UV light- and Fenton Reaction-induced oxidative DNA damage by the soybean isoflavone genistein. <i>Carcinogenesis</i> , 1996, 17, 73-77.	1.3	155
31	The relation of magnesium and calcium intakes and a genetic polymorphism in the magnesium transporter to colorectal neoplasia risk. <i>American Journal of Clinical Nutrition</i> , 2007, 86, 743-751.	2.2	155
32	Human metabolic correlates of body mass index. <i>Metabolomics</i> , 2014, 10, 259-269.	1.4	148
33	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. <i>Carcinogenesis</i> , 2014, 35, 1012-1019.	1.3	145
34	Metabolomics in Epidemiology: Sources of Variability in Metabolite Measurements and Implications. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 631-640.	1.1	144
35	Association of Genetic Polymorphisms in the VEGF Gene with Breast Cancer Survival. <i>Cancer Research</i> , 2005, 65, 5015-5019.	0.4	143
36	FTO genetic variants, dietary intake and body mass index: insights from 177 330 individuals. <i>Human Molecular Genetics</i> , 2014, 23, 6961-6972.	1.4	143

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37	A Common Deletion in the APOBEC3 Genes and Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2013, 105, 573-579.	3.0	141
38	Genome-Wide Association Study in East Asians Identifies Novel Susceptibility Loci for Breast Cancer. <i>PLoS Genetics</i> , 2012, 8, e1002532.	1.5	137
39	Genome-wide association analysis in East Asians identifies breast cancer susceptibility loci at 1q32.1, 5q14.3 and 15q26.1. <i>Nature Genetics</i> , 2014, 46, 886-890.	9.4	135
40	Association analyses of East Asian individuals and trans-ancestry analyses with European individuals reveal new loci associated with cholesterol and triglyceride levels. <i>Human Molecular Genetics</i> , 2017, 26, 1770-1784.	1.4	135
41	Exome sequencing generates high quality data in non-target regions. <i>BMC Genomics</i> , 2012, 13, 194.	1.2	130
42	Genomic Characterization of Esophageal Squamous Cell Carcinoma Reveals Critical Genes Underlying Tumorigenesis and Poor Prognosis. <i>American Journal of Human Genetics</i> , 2016, 98, 709-727.	2.6	129
43	Exome chip meta-analysis identifies novel loci and East Asian-specific coding variants that contribute to lipid levels and coronary artery disease. <i>Nature Genetics</i> , 2017, 49, 1722-1730.	9.4	129
44	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
45	Differential pre-malignant programs and microenvironment chart distinct paths to malignancy in human colorectal polyps. <i>Cell</i> , 2021, 184, 6262-6280.e26.	13.5	125
46	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
47	Telomere Length in White Blood Cell DNA and Lung Cancer: A Pooled Analysis of Three Prospective Cohorts. <i>Cancer Research</i> , 2014, 74, 4090-4098.	0.4	112
48	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , 2019, 156, 1455-1466.	0.6	111
49	Genetic polymorphism in the manganese superoxide dismutase gene, antioxidant intake, and breast cancer risk: results from the Shanghai Breast Cancer Study. <i>Breast Cancer Research</i> , 2004, 6, R647-55.	2.2	109
50	Identification of a Functional Genetic Variant at 16q12.1 for Breast Cancer Risk: Results from the Asia Breast Cancer Consortium. <i>PLoS Genetics</i> , 2010, 6, e1001002.	1.5	107
51	Longer Telomere Length in Peripheral White Blood Cells Is Associated with Risk of Lung Cancer and the rs2736100 (CLPTM1L-TERT) Polymorphism in a Prospective Cohort Study among Women in China. <i>PLoS ONE</i> , 2013, 8, e59230.	1.1	106
52	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
53	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
54	Common Variation in Vitamin D Pathway Genes Predicts Circulating 25-Hydroxyvitamin D Levels among African Americans. <i>PLoS ONE</i> , 2011, 6, e28623.	1.1	103

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55	Oxidative Stress, Obesity, and Breast Cancer Risk: Results From the Shanghai Women's Health Study. <i>Journal of Clinical Oncology</i> , 2009, 27, 2482-2488.	0.8	99
56	Circulating 25-Hydroxyvitamin D and Risk of Kidney Cancer: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. <i>American Journal of Epidemiology</i> , 2010, 172, 47-57.	1.6	98
57	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.	2.6	98
58	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016, 150, 1633-1645.	0.6	97
59	Soyfood intake and breast cancer survival: a followup of the Shanghai Breast Cancer Study. <i>Breast Cancer Research and Treatment</i> , 2005, 92, 11-17.	1.1	94
60	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. <i>PLoS ONE</i> , 2018, 13, e0198166.	1.1	94
61	Inhibition of 11 β -hydroxysteroid dehydrogenase type II selectively blocks the tumor COX-2 pathway and suppresses colon carcinogenesis in mice and humans. <i>Journal of Clinical Investigation</i> , 2009, 119, 876-885.	3.9	93
62	Distinct distribution and prognostic significance of molecular subtypes of breast cancer in Chinese women: a population-based cohort study. <i>BMC Cancer</i> , 2011, 11, 292.	1.1	93
63	Evaluation of Breast Cancer Susceptibility Loci in Chinese Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2357-2365.	1.1	92
64	Genome-wide association study identifies breast cancer risk variant at 10q21.2: results from the Asia Breast Cancer Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 4991-4999.	1.4	92
65	Genetic Polymorphisms in the TGF- β 1 Gene and Breast Cancer Survival. <i>Cancer Research</i> , 2004, 64, 836-839.	0.4	90
66	Drinking Green Tea Modestly Reduces Breast Cancer Risk. <i>Journal of Nutrition</i> , 2009, 139, 310-316.	1.3	90
67	Genetic and Clinical Predictors for Breast Cancer Risk Assessment and Stratification Among Chinese Women. <i>Journal of the National Cancer Institute</i> , 2010, 102, 972-981.	3.0	90
68	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
69	Predictors and Variability of Repeat Measurements of Urinary Phenols and Parabens in a Cohort of Shanghai Women and Men. <i>Environmental Health Perspectives</i> , 2014, 122, 733-740.	2.8	89
70	Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. <i>Human Molecular Genetics</i> , 2013, 22, 2539-2550.	1.4	86
71	APOBEC3 deletion polymorphism is associated with breast cancer risk among women of European ancestry. <i>Carcinogenesis</i> , 2013, 34, 2240-2243.	1.3	85
72	Singlet Oxygen Involvement in Ultraviolet (254 nm) Radiation-Induced Formation of 8-Hydroxy-Deoxyguanosine in DNA. <i>Free Radical Biology and Medicine</i> , 1997, 23, 148-154.	1.3	82

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73	Preclinical pharmacology of the natural product anticancer agent 10-hydroxycamptothecin, an inhibitor of topoisomerase I. <i>Cancer Chemotherapy and Pharmacology</i> , 1998, 41, 257-267.	1.1	81
74	Prospective study of oral microbiome and colorectal cancer risk in low-income and African American populations. <i>International Journal of Cancer</i> , 2019, 144, 2381-2389.	2.3	81
75	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 795-806.	0.9	81
76	Cruciferous vegetables, the GSTP1 IleVal genetic polymorphism, and breast cancer risk. <i>American Journal of Clinical Nutrition</i> , 2008, 87, 753-760.	2.2	80
77	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
78	Race, African Ancestry, and Helicobacter pylori Infection in a Low-Income United States Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 826-834.	1.1	76
79	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. <i>American Journal of Human Genetics</i> , 2015, 96, 5-20.	2.6	76
80	Evaluation of 11 Breast Cancer Susceptibility Loci in African-American Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2761-2764.	1.1	73
81	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
82	A Comprehensive cis-eQTL Analysis Revealed Target Genes in Breast Cancer Susceptibility Loci Identified in Genome-wide Association Studies. <i>American Journal of Human Genetics</i> , 2018, 102, 890-903.	2.6	72
83	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
84	A genome-wide association study of breast cancer in women of African ancestry. <i>Human Genetics</i> , 2013, 132, 39-48.	1.8	70
85	Genetic polymorphisms in uridine diphospho-glucuronosyltransferase 1A1 (UGT1A1) and risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2004, 85, 239-245.	1.1	68
86	Major metabolite of F2-isoprostane in urine may be a more sensitive biomarker of oxidative stress than isoprostane itself. <i>American Journal of Clinical Nutrition</i> , 2012, 96, 405-414.	2.2	68
87	A meta-analysis of genome-wide association studies for adiponectin levels in East Asians identifies a novel locus near WDR11-FGFR2. <i>Human Molecular Genetics</i> , 2014, 23, 1108-1119.	1.4	68
88	Home kitchen ventilation, cooking fuels, and lung cancer risk in a prospective cohort of never smoking women in Shanghai, China. <i>International Journal of Cancer</i> , 2015, 136, 632-638.	2.3	68
89	Endogenous Estrogens, Estrogen Metabolites, and Breast Cancer Risk in Postmenopausal Chinese Women. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw103.	3.0	67
90	Evaluating Genome-Wide Association Study-Identified Breast Cancer Risk Variants in African-American Women. <i>PLoS ONE</i> , 2013, 8, e58350.	1.1	66

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91	Association of Obesity-related Genetic Variants With Endometrial Cancer Risk: A Report From the Shanghai Endometrial Cancer Genetics Study. <i>American Journal of Epidemiology</i> , 2011, 174, 1115-1126.	1.6	65
92	Very Low-Level Heteroplasmy mtDNA Variations Are Inherited in Humans. <i>Journal of Genetics and Genomics</i> , 2013, 40, 607-615.	1.7	63
93	Genome-Wide Association Study Meta-Analysis Reveals Transethnic Replication of Mean Arterial and Pulse Pressure Loci. <i>Hypertension</i> , 2013, 62, 853-859.	1.3	63
94	Novel Genetic Markers of Breast Cancer Survival Identified by a Genome-Wide Association Study. <i>Cancer Research</i> , 2012, 72, 1182-1189.	0.4	62
95	Association of Leukocyte Telomere Length With Breast Cancer Risk: Nested Case-Control Findings From the Shanghai Women's Health Study. <i>American Journal of Epidemiology</i> , 2013, 177, 617-624.	1.6	62
96	Multiple Nonglycemic Genomic Loci Are Newly Associated With Blood Level of Glycated Hemoglobin in East Asians. <i>Diabetes</i> , 2014, 63, 2551-2562.	0.3	61
97	Genome-Wide Association Meta-analysis Identifies Novel Variants Associated With Fasting Plasma Glucose in East Asians. <i>Diabetes</i> , 2015, 64, 291-298.	0.3	59
98	Tumor tissue microRNA expression in association with triple-negative breast cancer outcomes. <i>Breast Cancer Research and Treatment</i> , 2015, 152, 183-191.	1.1	59
99	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. <i>American Journal of Human Genetics</i> , 2016, 99, 903-911.	2.6	59
100	Genetic polymorphisms in the estrogen receptor alpha gene and risk of breast cancer: results from the Shanghai Breast Cancer Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2003, 12, 853-9.	1.1	59
101	Dietary Folate Intake, MTHFR Genetic Polymorphisms, and the Risk of Endometrial Cancer among Chinese Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 281-287.	1.1	58
102	Genome-wide association studies in East Asians identify new loci for waist-hip ratio and waist circumference. <i>Scientific Reports</i> , 2016, 6, 17958.	1.6	58
103	Is high vitamin B12 status a cause of lung cancer?. <i>International Journal of Cancer</i> , 2019, 145, 1499-1503.	2.3	58
104	The Circadian Rhythm Gene <i>Arntl2</i> Is a Metastasis Susceptibility Gene for Estrogen Receptor-Negative Breast Cancer. <i>PLoS Genetics</i> , 2016, 12, e1006267.	1.5	57
105	Blood Vitamin D Levels in Relation to Genetic Estimation of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2325-2331.	1.1	56
106	Racial differences in the association between body mass index and serum IGF1, IGF2, and IGFBP3. <i>Endocrine-Related Cancer</i> , 2010, 17, 51-60.	1.6	56
107	Prospective Study of <i>Helicobacter pylori</i> Biomarkers for Gastric Cancer Risk among Chinese Men. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 2185-2192.	1.1	56
108	Urinary Levels of Trimethylamine-N-oxide and Incident Coronary Heart Disease: A Prospective Investigation Among Urban Chinese Adults. <i>Journal of the American Heart Association</i> , 2019, 8, e010606.	1.6	56

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109	Assessment of Dietary Isoflavone Intake among Middle-Aged Chinese Men1. <i>Journal of Nutrition</i> , 2007, 137, 1011-1016.	1.3	53
110	Association of Leukocyte Telomere Length with Colorectal Cancer Risk: Nested Caseâ€“Control Findings from the Shanghai Women's Health Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1807-1813.	1.1	53
111	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	1.4	53
112	<i>Helicobacter pylori</i> blood biomarker for gastric cancer risk in East Asia. <i>International Journal of Epidemiology</i> , 2016, 45, 774-781.	0.9	53
113	Dietary intake of PUFAs and colorectal polyp risk. <i>American Journal of Clinical Nutrition</i> , 2012, 95, 703-712.	2.2	52
114	Genomeâ€“wide association study identifies a new SMAD7 risk variant associated with colorectal cancer risk in East Asians. <i>International Journal of Cancer</i> , 2014, 135, 948-955.	2.3	52
115	Prediction of breast cancer risk based on common genetic variants in women of East Asian ancestry. <i>Breast Cancer Research</i> , 2016, 18, 124.	2.2	52
116	Isothiocyanate exposure, glutathione S-transferase polymorphisms, and colorectal cancer risk. <i>American Journal of Clinical Nutrition</i> , 2010, 91, 704-711.	2.2	51
117	Fineâ€“scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.	2.3	51
118	Variation in oral microbiome is associated with future risk of lung cancer among never-smokers. <i>Thorax</i> , 2021, 76, 256-263.	2.7	51
119	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
120	Genome-wide association studies in women of African ancestry identified 3q26.21 as a novel susceptibility locus for oestrogen receptor negative breast cancer. <i>Human Molecular Genetics</i> , 2016, 25, ddw305.	1.4	50
121	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
122	Quantitative analysis of mitochondrial DNA 4977-bp deletion in sporadic breast cancer and benign breast diseases. <i>Breast Cancer Research and Treatment</i> , 2008, 108, 427-434.	1.1	49
123	Biochemical Validation of Food Frequency Questionnaire-Estimated Carotenoid, Â-Tocopherol, and Folate Intakes Among African Americans and Non-Hispanic Whites in the Southern Community Cohort Study. <i>American Journal of Epidemiology</i> , 2010, 171, 488-497.	1.6	49
124	Urinary isoflavonoids and risk of coronary heart disease. <i>International Journal of Epidemiology</i> , 2012, 41, 1367-1375.	0.9	49
125	Genome-wide association study confirms lung cancer susceptibility loci on chromosomes 5p15 and 15q25 in an African-American population. <i>Lung Cancer</i> , 2016, 98, 33-42.	0.9	49
126	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2019, 79, 505-517.	0.4	49

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127	Age at menarche and age at natural menopause in East Asian women: a genome-wide association study. <i>Age</i> , 2016, 38, 513-523.	3.0	47
128	Integrative genomic analyses of APOBEC-mutational signature, expression and germline deletion of APOBEC3 genes, and immunogenicity in multiple cancer types. <i>BMC Medical Genomics</i> , 2019, 12, 131.	0.7	47
129	Oral microbiome and obesity in a large study of low-income and African-American populations. <i>Journal of Oral Microbiology</i> , 2019, 11, 1650597.	1.2	46
130	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.	5.8	46
131	Intra-Person Variation of Urinary Biomarkers of Oxidative Stress and Inflammation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 947-952.	1.1	45
132	Serum Adiponectin in Relation to Body Mass Index and Other Correlates in Black and White Women. <i>Annals of Epidemiology</i> , 2011, 21, 86-94.	0.9	45
133	Urinary Prostaglandin E2 Metabolite and Risk for Colorectal Adenoma. <i>Cancer Prevention Research</i> , 2012, 5, 336-342.	0.7	45
134	Visceral adiposity and risk of coronary heart disease in relatively lean Chinese adults. <i>International Journal of Cardiology</i> , 2013, 168, 2141-2145.	0.8	45
135	<i>Helicobacter pylori</i> Proteinâ€Specific Antibodies and Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 1964-1974.	1.1	45
136	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	3.0	45
137	Evaluation of Genetic Susceptibility Loci for Obesity in Chinese Women. <i>American Journal of Epidemiology</i> , 2010, 172, 244-254.	1.6	44
138	Prevalence and Determinants of Hyperuricemia in Middle-Aged, Urban Chinese Men. <i>Metabolic Syndrome and Related Disorders</i> , 2010, 8, 263-270.	0.5	44
139	Urinary Metabolite Risk Biomarkers of Lung Cancer: A Prospective Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 978-986.	1.1	44
140	Effects of caloric restriction on age-related oxidative modifications of macromolecules and lymphocyte proliferation in rats. <i>Free Radical Biology and Medicine</i> , 1995, 19, 859-865.	1.3	43
141	Identification of Novel Susceptibility Loci and Genes for Prostate Cancer Risk: A Transcriptome-Wide Association Study in Over 140,000 European Descendants. <i>Cancer Research</i> , 2019, 79, 3192-3204.	0.4	43
142	HTR1B, ADIPOR1, PPARGC1A, and CYP19A1 and Obesity in a Cohort of Caucasians and African Americans: An Evaluation of Gene-Environment Interactions and Candidate Genes. <i>American Journal of Epidemiology</i> , 2012, 175, 11-21.	1.6	42
143	Cooking Coal Use and All-Cause and Cause-Specific Mortality in a Prospective Cohort Study of Women in Shanghai, China. <i>Environmental Health Perspectives</i> , 2016, 124, 1384-1389.	2.8	42
144	Long-term diet quality is associated with gut microbiome diversity and composition among urban Chinese adults. <i>American Journal of Clinical Nutrition</i> , 2021, 113, 684-694.	2.2	42

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145	CYP2A6 reduced activity gene variants confer reduction in lung cancer risk in African American smokers—findings from two independent populations. <i>Carcinogenesis</i> , 2015, 36, 99-103.	1.3	41
146	MiR-374a suppresses lung adenocarcinoma cell proliferation and invasion by targeting <i>TGFA</i> gene expression. <i>Carcinogenesis</i> , 2016, 37, 567-575.	1.3	41
147	Expression patterns of the ATM gene in mammary tissues and their associations with breast cancer survival. <i>Cancer</i> , 2007, 109, 1729-1735.	2.0	40
148	Circulating transforming growth factor- β -1 and breast cancer prognosis: results from the Shanghai Breast Cancer Study. <i>Breast Cancer Research and Treatment</i> , 2008, 112, 335-341.	1.1	40
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