

Qiuyin Cai

List of Publications by Citations

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372
papers

17,552
citations

66
h-index

117
g-index

397
ext. papers

20,884
ext. citations

7.4
avg, IF

5.57
L-index

#	Paper	IF	Citations
372	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013 , 45, 353-61, 361e1-2	36.3	813
371	Genome-wide trans-ancestry meta-analysis provides insight into the genetic architecture of type 2 diabetes susceptibility. <i>Nature Genetics</i> , 2014 , 46, 234-44	36.3	784
370	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017 , 551, 92-94	50.4	643
369	Genome-wide association study identifies a new breast cancer susceptibility locus at 6q25.1. <i>Nature Genetics</i> , 2009 , 41, 324-8	36.3	434
368	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, 371-84, 384e1-2	36.3	422
367	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013 , 45, 385-91, 391e1-2	36.3	413
366	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015 , 47, 373-80	36.3	406
365	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013 , 45, 392-8, 398e1-2	36.3	327
364	Meta-analysis identifies common variants associated with body mass index in east Asians. <i>Nature Genetics</i> , 2012 , 44, 307-11	36.3	301
363	The landscape of recombination in African Americans. <i>Nature</i> , 2011 , 476, 170-5	50.4	243
362	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012 , 44, 1330-5	36.3	237
361	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
360	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. <i>Nature Genetics</i> , 2014 , 46, 533-42	36.3	175
359	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , 2011 , 43, 570-3	36.3	171
358	Identification of new genetic risk variants for type 2 diabetes. <i>PLoS Genetics</i> , 2010 , 6, e1001127	6	168
357	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	11	167
356	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	9.9	165

355	Urinary isothiocyanate levels, brassica, and human breast cancer. <i>Cancer Research</i> , 2003 , 63, 3980-6	10.1	158
354	Genome-wide association analyses in East Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2013 , 45, 191-6	36.3	155
353	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	3.8	155
352	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	6	151
351	Inhibition of UV light- and Fenton reaction-induced oxidative DNA damage by the soybean isoflavone genistein. <i>Carcinogenesis</i> , 1996 , 17, 73-7	4.6	147
350	Southern community cohort study: establishing a cohort to investigate health disparities. <i>Journal of the National Medical Association</i> , 2005 , 97, 972-9	2.3	144
349	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, 5492-504	5.6	141
348	The 5p15.33 locus is associated with risk of lung adenocarcinoma in never-smoking females in Asia. <i>PLoS Genetics</i> , 2010 , 6, e1001051	6	141
347	The Human Tumor Atlas Network: Charting Tumor Transitions across Space and Time at Single-Cell Resolution. <i>Cell</i> , 2020 , 181, 236-249	56.2	140
346	Association of genetic polymorphisms in the VEGF gene with breast cancer survival. <i>Cancer Research</i> , 2005 , 65, 5015-9	10.1	133
345	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, 743-51	7	125
344	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. <i>Carcinogenesis</i> , 2014 , 35, 1012-9	4.6	121
343	FTO genetic variants, dietary intake and body mass index: insights from 177,330 individuals. <i>Human Molecular Genetics</i> , 2014 , 23, 6961-72	5.6	120
342	Genome-wide association study in east Asians identifies novel susceptibility loci for breast cancer. <i>PLoS Genetics</i> , 2012 , 8, e1002532	6	118
341	Human metabolic correlates of body mass index. <i>Metabolomics</i> , 2014 , 10, 259-269	4.7	111
340	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, 886-90	36.3	110
339	A common deletion in the APOBEC3 genes and breast cancer risk. <i>Journal of the National Cancer Institute</i> , 2013 , 105, 573-9	9.7	109
338	Metabolomics in epidemiology: sources of variability in metabolite measurements and implications. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 631-40	4	109

337	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, 1052-67	24.4	104
336	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	36.3	101
335	Exome sequencing generates high quality data in non-target regions. <i>BMC Genomics</i> , 2012 , 13, 194	4.5	95
334	Genomic Characterization of Esophageal Squamous Cell Carcinoma Reveals Critical Genes Underlying Tumorigenesis and Poor Prognosis. <i>American Journal of Human Genetics</i> , 2016 , 98, 709-27	11	95
333	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016 , 48, 374-86	36.3	93
332	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	6	93
331	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	8.3	93
330	Common variation in vitamin D pathway genes predicts circulating 25-hydroxyvitamin D Levels among African Americans. <i>PLoS ONE</i> , 2011 , 6, e28623	3.7	92
329	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	5.6	90
328	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	3.7	90
327	Telomere length in white blood cell DNA and lung cancer: a pooled analysis of three prospective cohorts. <i>Cancer Research</i> , 2014 , 74, 4090-8	10.1	88
326	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014 , 4, 4999	17.4	87
325	Evaluation of breast cancer susceptibility loci in Chinese women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 2357-65	4	87
324	Genetic polymorphisms in the TGF-beta 1 gene and breast cancer survival: a report from the Shanghai Breast Cancer Study. <i>Cancer Research</i> , 2004 , 64, 836-9	10.1	87
323	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	36.3	83
322	Oxidative stress, obesity, and breast cancer risk: results from the Shanghai Women@ Health Study. <i>Journal of Clinical Oncology</i> , 2009 , 27, 2482-8	2.2	83
321	Inhibition of 11beta-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, 876-85	15.9	82
320	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, 1046-60	11	80

319	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	4.8	80
318	Soyfood intake and breast cancer survival: a followup of the Shanghai Breast Cancer Study. <i>Breast Cancer Research and Treatment</i> , 2005 , 92, 11-7	4.4	80
317	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, 4991-9	5.6	79
316	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, 6616-33	5.6	77
315	Drinking green tea modestly reduces breast cancer risk. <i>Journal of Nutrition</i> , 2009 , 139, 310-6	4.1	77
314	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, 2539-50	5.6	75
313	APOBEC3 deletion polymorphism is associated with breast cancer risk among women of European ancestry. <i>Carcinogenesis</i> , 2013 , 34, 2240-3	4.6	74
312	Genetic and clinical predictors for breast cancer risk assessment and stratification among Chinese women. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 972-81	9.7	73
311	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, 733-40	8.4	72
310	Cruciferous vegetables, the GSTP1 Ile105Val genetic polymorphism, and breast cancer risk. <i>American Journal of Clinical Nutrition</i> , 2008 , 87, 753-60	7	72
309	Meta-analysis of genome-wide association studies of adult height in East Asians identifies 17 novel loci. <i>Human Molecular Genetics</i> , 2015 , 24, 1791-800	5.6	71
308	Evaluation of 11 breast cancer susceptibility loci in African-American women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 2761-4	4	71
307	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, 148-54	7.8	66
306	Preclinical pharmacology of the natural product anticancer agent 10-hydroxycamptothecin, an inhibitor of topoisomerase I. <i>Cancer Chemotherapy and Pharmacology</i> , 1998 , 41, 257-67	3.5	66
305	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016 , 150, 1633-1645	13.3	64
304	A genome-wide association study of breast cancer in women of African ancestry. <i>Human Genetics</i> , 2013 , 132, 39-48	6.3	63
303	Race, African ancestry, and Helicobacter pylori infection in a low-income United States population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011 , 20, 826-34	4	61
302	Genetic polymorphisms in uridine diphospho-glucuronosyltransferase 1A1 (UGT1A1) and risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2004 , 85, 239-45	4.4	61

301	Genome-wide association study meta-analysis reveals transethnic replication of mean arterial and pulse pressure loci. <i>Hypertension</i> , 2013 , 62, 853-9	8.5	60
300	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, 1344-55	10.1	60
299	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	11	59
298	Evaluating genome-wide association study-identified breast cancer risk variants in African-American women. <i>PLoS ONE</i> , 2013 , 8, e58350	3.7	58
297	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	4	58
296	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, 405-14	7	56
295	Novel genetic markers of breast cancer survival identified by a genome-wide association study. <i>Cancer Research</i> , 2012 , 72, 1182-9	10.1	56
294	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020 , 52, 56-73	36.3	56
293	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, 311-9	7.5	55
292	Dietary folate intake, MTHFR genetic polymorphisms, and the risk of endometrial cancer among Chinese women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 281-7	4	55
291	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , 2019 , 156, 1455-1466	13.3	55
290	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016 , 7, 12675	17.4	53
289	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, 1108-19	5.6	53
288	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, 617-24	3.8	52
287	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	7.8	52
286	Prospective study of Helicobacter pylori biomarkers for gastric cancer risk among Chinese men. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 2185-92	4	50
285	Tumor tissue microRNA expression in association with triple-negative breast cancer outcomes. <i>Breast Cancer Research and Treatment</i> , 2015 , 152, 183-191	4.4	49
284	Genome-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, 948-55	7.5	49

283	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, 1115-26	3.8	49
282	Association of leukocyte telomere length with colorectal cancer risk: nested case-control findings from the Shanghai Women's Health Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 1807-13	4	49
281	Genome-wide association studies in East Asians identify new loci for waist-hip ratio and waist circumference. <i>Scientific Reports</i> , 2016 , 6, 17958	4.9	48
280	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014 , 23, 6096-111	5.6	48
279	Multiple nonglycemic genomic loci are newly associated with blood level of glycated hemoglobin in East Asians. <i>Diabetes</i> , 2014 , 63, 2551-62	0.9	46
278	Blood vitamin d levels in relation to genetic estimation of African ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 2325-31	4	46
277	Endogenous Estrogens, Estrogen Metabolites, and Breast Cancer Risk in Postmenopausal Chinese Women. <i>Journal of the National Cancer Institute</i> , 2016 , 108,	9.7	46
276	Very low-level heteroplasmy mtDNA variations are inherited in humans. <i>Journal of Genetics and Genomics</i> , 2013 , 40, 607-15	4	45
275	Isothiocyanate exposure, glutathione S-transferase polymorphisms, and colorectal cancer risk. <i>American Journal of Clinical Nutrition</i> , 2010 , 91, 704-11	7	45
274	Racial differences in the association between body mass index and serum IGF1, IGF2, and IGFBP3. <i>Endocrine-Related Cancer</i> , 2010 , 17, 51-60	5.7	45
273	Assessment of dietary isoflavone intake among middle-aged Chinese men. <i>Journal of Nutrition</i> , 2007 , 137, 1011-6	4.1	45
272	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, 427-34	4.4	45
271	Biochemical validation of food frequency questionnaire-estimated carotenoid, alpha-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, 488-97	3.8	44
270	The trans-ancestral genomic architecture of glycemic traits. <i>Nature Genetics</i> , 2021 , 53, 840-860	36.3	44
269	Genome-wide association meta-analysis identifies novel variants associated with fasting plasma glucose in East Asians. <i>Diabetes</i> , 2015 , 64, 291-8	0.9	43
268	Dietary intake of PUFAs and colorectal polyp risk. <i>American Journal of Clinical Nutrition</i> , 2012 , 95, 703-127		43
267	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	11	43
266	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	7.5	43

265	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 ¹¹	42
264	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, 632-8	41
263	Urinary isoflavonoids and risk of coronary heart disease. <i>International Journal of Epidemiology</i> , 2012 , 41, 1367-75	41
262	Evaluation of genetic susceptibility loci for obesity in Chinese women. <i>American Journal of Epidemiology</i> , 2010 , 172, 244-54	40
261	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> , 2017 , 26, 454-465	40
260	Prevalence and determinants of hyperuricemia in middle-aged, urban Chinese men. <i>Metabolic Syndrome and Related Disorders</i> , 2010 , 8, 263-70	39
259	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	38
258	Helicobacter pylori protein-specific antibodies and risk of colorectal cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 1964-74	37
257	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	37
256	Urinary prostaglandin E2 metabolite and risk for colorectal adenoma. <i>Cancer Prevention Research</i> , 2012 , 5, 336-42	37
255	Common MMP-7 polymorphisms and breast cancer susceptibility: a multistage study of association and functionality. <i>Cancer Research</i> , 2008 , 68, 6453-9	37
254	Expression patterns of the ATM gene in mammary tissues and their associations with breast cancer survival. <i>Cancer</i> , 2007 , 109, 1729-35	37
253	Insulin-like growth factor-I gene polymorphism and breast cancer risk in Chinese women. <i>International Journal of Cancer</i> , 2005 , 113, 307-11	37
252	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, 859-65	37
251	Helicobacter pylori blood biomarker for gastric cancer risk in East Asia. <i>International Journal of Epidemiology</i> , 2016 , 45, 774-81	37
250	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015 , 24, 2966-84	36
249	Intra-person variation of urinary biomarkers of oxidative stress and inflammation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 947-52	36
248	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015 , 24, 285-98	35

247	Genetic variant in TP63 on locus 3q28 is associated with risk of lung adenocarcinoma among never-smoking females in Asia. <i>Human Genetics</i> , 2012 , 131, 1197-203	6.3	35
246	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	3.8	35
245	Immunohistochemical expressions of Ki-67, cyclin D1, beta-catenin, cyclooxygenase-2, and epidermal growth factor receptor in human colorectal adenoma: a validation study of tissue microarrays. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1719-26	4	35
244	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	6	35
243	Circulating transforming growth factor-beta-1 and breast cancer prognosis: results from the Shanghai Breast Cancer Study. <i>Breast Cancer Research and Treatment</i> , 2008 , 112, 335-41	4.4	34
242	Prediction of breast cancer risk based on common genetic variants in women of East Asian ancestry. <i>Breast Cancer Research</i> , 2016 , 18, 124	8.3	34
241	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, 4835-4846	5.6	34
240	Joint effect of estrogen receptor beta sequence variants and endogenous estrogen exposure on breast cancer risk in Chinese women. <i>Cancer Research</i> , 2003 , 63, 7624-9	10.1	34
239	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	4.6	33
238	Genetic polymorphisms in the MMP-7 gene and breast cancer survival. <i>International Journal of Cancer</i> , 2009 , 124, 208-14	7.5	33
237	Is high vitamin B12 status a cause of lung cancer?. <i>International Journal of Cancer</i> , 2019 , 145, 1499-1503	7.5	33
236	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, 620-9	5.6	32
235	Plasma carotenoids, tocopherols, retinol and breast cancer risk: results from the Shanghai Women Health Study (SWHS). <i>Breast Cancer Research and Treatment</i> , 2009 , 117, 381-9	4.4	32
234	Evaluation of genome-wide association study-identified type 2 diabetes loci in African Americans. <i>American Journal of Epidemiology</i> , 2012 , 176, 995-1001	3.8	32
233	MiR-374a suppresses lung adenocarcinoma cell proliferation and invasion by targeting TGFA gene expression. <i>Carcinogenesis</i> , 2016 , 37, 567-75	4.6	32
232	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	3.7	31
231	Circulating C-reactive protein and colorectal cancer risk: a report from the Shanghai MenQ Health Study. <i>Carcinogenesis</i> , 2013 , 34, 2799-803	4.6	31
230	Polymorphisms in tissue inhibitors of metalloproteinases-2 and -3 and breast cancer susceptibility and survival. <i>International Journal of Cancer</i> , 2009 , 125, 844-50	7.5	31

229	Common genetic variants in the vitamin D pathway including genome-wide associated variants are not associated with breast cancer risk among Chinese women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011 , 20, 2313-6	4	31
228	Polymorphisms in the CYP19A1 (aromatase) gene and endometrial cancer risk in Chinese women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 943-9	4	31
227	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	5.9	31
226	Circulating cytokines and gastric cancer risk. <i>Cancer Causes and Control</i> , 2013 , 24, 2245-50	2.8	30
225	Visceral adiposity and risk of coronary heart disease in relatively lean Chinese adults. <i>International Journal of Cardiology</i> , 2013 , 168, 2141-5	3.2	30
224	Circulating 25-hydroxyvitamin D levels in relation to blood pressure parameters and hypertension in the Shanghai WomenQ and MenQ Health Studies. <i>British Journal of Nutrition</i> , 2012 , 108, 449-58	3.6	30
223	Genome-wide association study identifies a possible susceptibility locus for endometrial cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 980-7	4	30
222	Polymorphisms and haplotypes in the caspase-3, caspase-7, and caspase-8 genes and risk for endometrial cancer: a population-based, case-control study in a Chinese population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 2114-22	4	30
221	Urinary polyphenols and breast cancer risk: results from the Shanghai WomenQ Health Study. <i>Breast Cancer Research and Treatment</i> , 2010 , 120, 693-702	4.4	30
220	Genetic polymorphisms of metastasis suppressor gene NME1 and breast cancer survival. <i>Clinical Cancer Research</i> , 2008 , 14, 4787-93	12.9	30
219	The Circadian Rhythm Gene Arntl2 Is a Metastasis Susceptibility Gene for Estrogen Receptor-Negative Breast Cancer. <i>PLoS Genetics</i> , 2016 , 12, e1006267	6	30
218	Age at menarche and age at natural menopause in East Asian women: a genome-wide association study. <i>Age</i> , 2016 , 38, 513-523		30
217	Dietary B vitamin and methionine intakes and plasma folate are not associated with colorectal cancer risk in Chinese women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 1003-6	4	29
216	Promoter methylation status of the MGMT, hMLH1, and CDKN2A/p16 genes in non-neoplastic mucosa of patients with and without colorectal adenomas. <i>Oncology Reports</i> , 2006 , 16, 429-35	3.5	29
215	Dietary isoflavones, urinary isoflavonoids, and risk of ischemic stroke in women. <i>American Journal of Clinical Nutrition</i> , 2015 , 102, 680-6	7	28
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