Anna H Wu

List of Publications by Year in descending order

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285 papers 18,674 citations

64 h-index 121 g-index

288 all docs

288 docs citations

times ranked

288

20561 citing authors

#	Article	IF	Citations
1	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
3	Association between endometriosis and risk of histological subtypes of ovarian cancer: a pooled analysis of case–control studies. Lancet Oncology, The, 2012, 13, 385-394.	5.1	753
4	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 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. Nature Genetics, 2013, 45, 371-384.	9.4	493
6	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377
7	Genome-wide association studies identify four ER negative–specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	9.4	374
8	Adolescent and adult soy intake and risk of breast cancer in Asian-Americans. Carcinogenesis, 2002, 23, 1491-1496.	1.3	359
9	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	9.4	356
10	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 2013, 45, 362-370.	9.4	326
11	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. Nature Genetics, 2010, 42, 874-879.	9.4	321
12	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
13	Meta-analysis: Dietary Fat Intake, Serum Estrogen Levels, and the Risk of Breast Cancer. Journal of the National Cancer Institute, 1999, 91, 529-534.	3.0	283
14	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. Nature Genetics, 2009, 41, 996-1000.	9.4	276
15	A multiethnic population-based study of smoking, alcohol and body size and risk of adenocarcinomas of the stomach and esophagus (United States). Cancer Causes and Control, 2001, 12, 721-732.	0.8	264
16	Body mass index in relation to oesophageal and oesophagogastric junction adenocarcinomas: a pooled analysis from the International BEACON Consortium. International Journal of Epidemiology, 2012, 41, 1706-1718.	0.9	237
17	Common variants at 19p13 are associated with susceptibility to ovarian cancer. Nature Genetics, 2010, 42, 880-884.	9.4	235
18	Green tea and risk of breast cancer in Asian Americans. International Journal of Cancer, 2003, 106, 574-579.	2.3	226

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19	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	9.4	221
20	Hiatal hernia, reflux symptoms, body size, and risk of esophageal and gastric adenocarcinoma. Cancer, 2003, 98, 940-948.	2.0	204
21	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. Nature Communications, 2020, 11, 597.	5.8	193
22	Aspirin, Nonaspirin Nonsteroidal Anti-inflammatory Drug, and Acetaminophen Use and Risk of Invasive Epithelial Ovarian Cancer: A Pooled Analysis in the Ovarian Cancer Association Consortium. Journal of the National Cancer Institute, 2014, 106, djt431-djt431.	3.0	186
23	A genome-wide association study identifies new susceptibility loci for esophageal adenocarcinoma and Barrett's esophagus. Nature Genetics, 2013, 45, 1487-1493.	9.4	174
24	Obesity and risk of ovarian cancer subtypes: evidence from the Ovarian Cancer Association Consortium. Endocrine-Related Cancer, 2013, 20, 251-262.	1.6	169
25	Role ofHelicobacter pylori CagA+ strains and risk of adenocarcinoma of the stomach and esophagus. International Journal of Cancer, 2003, 103, 815-821.	2.3	162
26	Common variants at the MHC locus and at chromosome 16q24.1 predispose to Barrett's esophagus. Nature Genetics, 2012, 44, 1131-1136.	9.4	162
27	Mammographic density and breast cancer in three ethnic groups. Cancer Epidemiology Biomarkers and Prevention, 2003, 12, 332-8.	1.1	158
28	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	7.7	157
29	Dietary patterns and breast cancer risk in Asian American women. American Journal of Clinical Nutrition, 2009, 89, 1145-1154.	2.2	148
30	Tubal ligation and risk of ovarian cancer subtypes: a pooled analysis of case-control studies. International Journal of Epidemiology, 2013, 42, 579-589.	0.9	146
31	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	5.8	144
32	Tea intake, COMT genotype, and breast cancer in Asian-American women. Cancer Research, 2003, 63, 7526-9.	0.4	143
33	Gastroesophageal Reflux in Relation to Adenocarcinomas of the Esophagus: A Pooled Analysis from the Barrett's and Esophageal Adenocarcinoma Consortium (BEACON). PLoS ONE, 2014, 9, e103508.	1.1	134
34	Genome-wide association studies in oesophageal adenocarcinoma and Barrett's oesophagus: a large-scale meta-analysis. Lancet Oncology, The, 2016, 17, 1363-1373.	5.1	133
35	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386.	9.4	125
36	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	2.6	124

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37	Hormonal factors and the risk of invasive ovarian cancer: a population-based case-control study. Fertility and Sterility, 2004, 82, 186-195.	0.5	122
38	Sleep duration, melatonin and breast cancer among Chinese women in Singapore. Carcinogenesis, 2008, 29, 1244-1248.	1.3	121
39	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
40	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 1619-1630.	0.9	111
41	Cumulative Burden of Colorectal Cancer–Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	0.6	110
42	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	5.8	105
43	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1156-1166.	1.1	101
44	Markers of inflammation and risk of ovarian cancer in Los Angeles County. International Journal of Cancer, 2009, 124, 1409-1415.	2.3	100
45	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. Nature Communications, 2013, 4, 1627.	5.8	98
46	Increased ovarian cancer risk associated with menopausal estrogen therapy is reduced by adding a progestin. Cancer, 2009, 115, 531-539.	2.0	97
47	A vegetable-fruit-soy dietary pattern protects against breast cancer among postmenopausal Singapore Chinese women. American Journal of Clinical Nutrition, 2010, 91, 1013-1019.	2.2	96
48	No evidence that protein truncating variants in <i>BRIP1</i> are associated with breast cancer risk: implications for gene panel testing. Journal of Medical Genetics, 2016, 53, 298-309.	1.5	94
49	Polymorphisms Near TBX5 and GDF7 Are Associated With Increased Risk for Barrett's Esophagus. Gastroenterology, 2015, 148, 367-378.	0.6	93
50	Effect of 2-Month Controlled Green Tea Intervention on Lipoprotein Cholesterol, Glucose, and Hormone Levels in Healthy Postmenopausal Women. Cancer Prevention Research, 2012, 5, 393-402.	0.7	91
51	Obesity and Mortality After Breast Cancer by Race/Ethnicity: The California Breast Cancer Survivorship Consortium. American Journal of Epidemiology, 2014, 179, 95-111.	1.6	90
52	Circulating Levels of Insulin-like Growth Factor 1 and Insulin-like Growth Factor Binding Protein 3 Associate With Risk of Colorectal Cancer Based on Serologic and Mendelian Randomization Analyses. Gastroenterology, 2020, 158, 1300-1312.e20.	0.6	90
53	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
54	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. Nature Communications, 2020, 11, 3833.	5.8	88

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55	Germline Genetic Contributions to Risk for Esophageal Adenocarcinoma, Barrett's Esophagus, and Gastroesophageal Reflux. Journal of the National Cancer Institute, 2013, 105, 1711-1718.	3.0	85
56	Effects of green tea catechin extract on serum lipids in postmenopausal women: a randomized, placebo-controlled clinical trial. American Journal of Clinical Nutrition, 2016, 104, 1671-1682.	2.2	85
57	Cigarette smoking and risk of ovarian cancer: a pooled analysis of 21 case–control studies. Cancer Causes and Control, 2013, 24, 989-1004.	0.8	84
58	Population Distribution of Lifetime Risk of Ovarian Cancer in the United States. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 671-676.	1.1	82
59	Smoking, alcohol use, dietary factors and risk of small intestinal adenocarcinoma. International Journal of Cancer, 1997, 70, 512-517.	2.3	79
60	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast–ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
61	Association Between Breastfeeding and Ovarian Cancer Risk. JAMA Oncology, 2020, 6, e200421.	3.4	78
62	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. American Journal of Human Genetics, 2015, 96, 5-20.	2.6	76
63	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. BMC Medicine, 2020, 18, 396.	2.3	76
64	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799.	0.4	75
65	Consortium analysis of 7 candidate SNPs for ovarian cancer. International Journal of Cancer, 2008, 123, 380-388.	2.3	73
66	Generalizability and Epidemiologic Characterization of Eleven Colorectal Cancer GWAS Hits in Multiple Populations. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 70-81.	1.1	73
67	Tea and circulating estrogen levels in postmenopausal Chinese women in Singapore. Carcinogenesis, 2005, 26, 976-980.	1.3	72
68	A Randomized Controlled Trial of Green Tea Extract Supplementation and Mammographic Density in Postmenopausal Women at Increased Risk of Breast Cancer. Cancer Prevention Research, 2017, 10, 710-718.	0.7	72
69	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 884-895.	0.9	71
70	The safety of green tea extract supplementation in postmenopausal women at risk for breast cancer: results of the Minnesota Green TeaÂTrial. Food and Chemical Toxicology, 2015, 83, 26-35.	1.8	69
71	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. Human Molecular Genetics, 2015, 24, 5955-5964.	1.4	68
72	Physical activity and breast cancer risk among Asian-American women in Los Angeles. Cancer, 2003, 97, 2565-2575.	2.0	67

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73	Determining Risk of Barrett's Esophagus and Esophageal Adenocarcinoma Based on Epidemiologic Factors and GeneticÂVariants. Gastroenterology, 2018, 154, 1273-1281.e3.	0.6	67
74	Fiber intake and risk of adenocarcinomas of the esophagus and stomach. Cancer Causes and Control, 2007, 18, 713-722.	0.8	66
75	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. Nature Communications, 2015, 6, 8234.	5.8	63
76	Dense breast stromal tissue shows greatly increased concentration of breast epithelium but no increase in its proliferative activity. Breast Cancer Research, 2006, 8, R24.	2.2	62
77	Diabetes and Other Comorbidities in Breast Cancer Survival by Race/Ethnicity: The California Breast Cancer Survivorship Consortium (CBCSC). Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 361-368.	1.1	62
78	Pelvic Inflammatory Disease and the Risk of Ovarian Cancer and Borderline Ovarian Tumors: A Pooled Analysis of 13 Case-Control Studies. American Journal of Epidemiology, 2017, 185, 8-20.	1.6	61
79	Association between ambient air pollution and breast cancer risk: The multiethnic cohort study. International Journal of Cancer, 2020, 146, 699-711.	2.3	60
80	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. American Journal of Human Genetics, 2016, 99, 903-911.	2.6	59
81	Body size, hormone therapy and risk of breast cancer in Asian-American women. International Journal of Cancer, 2007, 120, 844-852.	2.3	56
82	Reproductive history, breastâ€feeding and risk of triple negative breast cancer: The Breast Cancer Etiology in Minorities (BEM) study. International Journal of Cancer, 2018, 142, 2273-2285.	2.3	56
83	Plasma isoflavone levels versus self-reported soy isoflavone levels in Asian-American women in Los Angeles County. Carcinogenesis, 2003, 25, 77-81.	1.3	55
84	A promoter polymorphism in the CASP8 gene is not associated with cancer risk. Nature Genetics, 2008, 40, 259-260.	9.4	54
85	Combined and Interactive Effects of Environmental and GWAS-Identified Risk Factors in Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 880-890.	1.1	54
86	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. Cancer Research, 2018, 78, 5419-5430.	0.4	54
87	Evaluation of Medicare Claims Data as a Tool to Identify Dementia. Journal of Alzheimer's Disease, 2019, 67, 769-778.	1.2	54
88	Invasive breast cancer incidence trends by detailed race/ethnicity and age. International Journal of Cancer, 2012, 130, 395-404.	2.3	53
89	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2014, 23, 6096-6111.	1.4	53
90	Intersection of Race/Ethnicity and Socioeconomic Status in Mortality After Breast Cancer. Journal of Community Health, 2015, 40, 1287-1299.	1.9	53

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91	Nonsteroidal Anti-inflammatory Drugs and Risk of Esophageal and Gastric Adenocarcinomas in Los Angeles County. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 126-134.	1.1	52
92	Prediction of breast cancer risk based on common genetic variants in women of East Asian ancestry. Breast Cancer Research, 2016, 18, 124.	2.2	52
93	Green and black tea in relation to gynecologic cancers. Molecular Nutrition and Food Research, 2011, 55, 931-940.	1.5	51
94	Fineâ€scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. International Journal of Cancer, 2016, 139, 1303-1317.	2.3	51
95	Association of Dietary Vitamin A, Carotenoids, and Other Antioxidants with the Risk of Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 669-676.	1.1	50
96	Occupational physical activity and risk of adenocarcinomas of the esophagus and stomach. International Journal of Cancer, 2006, 118, 1004-1009.	2.3	49
97	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. Cancer Research, 2019, 79, 505-517.	0.4	49
98	Gut microbiome associations with breast cancer risk factors and tumor characteristics: a pilot study. Breast Cancer Research and Treatment, 2020, 182, 451-463.	1.1	48
99	A controlled 2-mo dietary fat reduction and soy food supplementation study in postmenopausal women. American Journal of Clinical Nutrition, 2005, 81, 1133-1141.	2.2	47
100	The California Breast Cancer Survivorship Consortium (CBCSC): prognostic factors associated with racial/ethnic differences in breast cancer survival. Cancer Causes and Control, 2013, 24, 1821-1836.	0.8	47
101	Tea, hormone-related cancers and endogenous hormone levels. Molecular Nutrition and Food Research, 2006, 50, 160-169.	1.5	46
102	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. Nature Communications, 2020, 11, 1217.	5.8	46
103	Sleep duration, spot urinary 6â€sulfatoxymelatonin levels and risk of breast cancer among Chinese women in Singapore. International Journal of Cancer, 2013, 132, 891-896.	2.3	45
104	Effect of Green Tea Supplements on Liver Enzyme Elevation: Results from a Randomized Intervention Study in the United States. Cancer Prevention Research, 2017, 10, 571-579.	0.7	45
105	Green tea and breast cancer. Molecular Nutrition and Food Research, 2011, 55, 921-930.	1.5	44
106	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. PLoS ONE, 2015, 10, e0128106.	1.1	44
107	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. Breast Cancer Research, 2017, 19, 119.	2.2	43
108	Development and Validation of the Gene Expression Predictor of High-grade Serous Ovarian Carcinoma Molecular SubTYPE (PrOTYPE). Clinical Cancer Research, 2020, 26, 5411-5423.	3.2	43

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109	Breast Cancer and Oral Contraceptive Use in Asian-American Women. American Journal of Epidemiology, 1999, 150, 561-567.	1.6	42
110	Comprehensive association testing of common genetic variation in DNA repair pathway genes in relationship with breast cancer risk in multiple populations. Human Molecular Genetics, 2008, 17, 825-834.	1.4	42
111	Diabetes and risk of breast cancer in Asian-American women. Carcinogenesis, 2007, 28, 1561-1566.	1.3	41
112	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	1.4	40
113	Association Between Menopausal Estrogen-Only Therapy and Ovarian Carcinoma Risk. Obstetrics and Gynecology, 2016, 127, 828-836.	1.2	39
114	Recreational physical inactivity and mortality in women with invasive epithelial ovarian cancer: evidence from the Ovarian Cancer Association Consortium. British Journal of Cancer, 2016, 115, 95-101.	2.9	39
115	The Minnesota Green Tea Trial (MGTT), a randomized controlled trial of the efficacy of green tea extract on biomarkers of breast cancer risk: study rationale, design, methods, and participant characteristics. Cancer Causes and Control, 2015, 26, 1405-1419.	0.8	38
116	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. Human Molecular Genetics, 2015, 24, 285-298.	1.4	38
117	Evaluating genetic variants associated with breast cancer risk in high and moderate-penetrance genes in Asians. Carcinogenesis, 2017, 38, 511-518.	1.3	38
118	Germline variation in inflammation-related pathways and risk of Barrett's oesophagus and oesophageal adenocarcinoma. Gut, 2017, 66, 1739-1747.	6.1	38
119	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. American Journal of Human Genetics, 2015, 97, 22-34.	2.6	37
120	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. International Journal of Epidemiology, 2019, 48, 781-794.	0.9	37
121	Associations Between Soy, Diet, Reproductive Factors, and Mammographic Density in Singapore Chinese Women. Nutrition and Cancer, 2006, 56, 128-135.	0.9	35
122	Green and black tea intake in relation to prostate cancer risk among Singapore Chinese. Cancer Causes and Control, 2012, 23, 1635-1641.	0.8	35
123	Associations Between Glycemic Traits and Colorectal Cancer: A Mendelian Randomization Analysis. Journal of the National Cancer Institute, 2022, 114, 740-752.	3.0	35
124	Tamoxifen, Soy, and Lifestyle Factors in Asian American Women With Breast Cancer. Journal of Clinical Oncology, 2007, 25, 3024-3030.	0.8	33
125	African Americans and Hispanics Remain at Lower Risk of Ovarian Cancer Than Non-Hispanic Whites after Considering Nongenetic Risk Factors and Oophorectomy Rates. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1094-1100.	1.1	33
126	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	3.2	33

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127	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. Human Molecular Genetics, 2016, 25, 3863-3876.	1.4	33
128	Racial/ethnic differences in the epidemiology of ovarian cancer: a pooled analysis of 12 case-control studies. International Journal of Epidemiology, 2018, 47, 460-472.	0.9	33
129	Urinary phthalate exposures and risk of breast cancer: the Multiethnic Cohort study. Breast Cancer Research, 2021, 23, 44.	2.2	33
130	Double-Blind Randomized 12-Month Soy Intervention Had No Effects on Breast MRI Fibroglandular Tissue Density or Mammographic Density. Cancer Prevention Research, 2015, 8, 942-951.	0.7	32
131	Chronic Recreational Physical Inactivity and Epithelial Ovarian Cancer Risk: Evidence from the Ovarian Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1114-1124.	1.1	32
132	Risk Prediction for Epithelial Ovarian Cancer in 11 United States–Based Case-Control Studies: Incorporation of Epidemiologic Risk Factors and 17 Confirmed Genetic Loci. American Journal of Epidemiology, 2016, 184, 555-569.	1.6	32
133	Body mass index, comorbidities, and hormonal factors in relation to meningioma in an ethnically diverse population: the Multiethnic Cohort. Neuro-Oncology, 2019, 21, 498-507.	0.6	32
134	Integrative post-genome-wide association analysis of CDKN2A and TP53 SNPs and risk of esophageal adenocarcinoma. Carcinogenesis, 2014, 35, 2740-2747.	1.3	31
135	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. Breast Cancer Research, 2016, 18, 64.	2.2	31
136	Age-specific risk factor profiles of adenocarcinomas of the esophagus: A pooled analysis from the international BEACON consortium. International Journal of Cancer, 2016, 138, 55-64.	2.3	31
137	Hormone therapy, DNA methylation and colon cancer. Carcinogenesis, 2010, 31, 1060-1067.	1.3	30
138	Green Tea, Soy, and Mammographic Density in Singapore Chinese Women. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 3358-3365.	1.1	29
139	Contribution of the Neighborhood Environment and Obesity to Breast Cancer Survival: The California Breast Cancer Survivorship Consortium. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1282-1290.	1.1	29
140	Network-Based Integration of GWAS and Gene Expression Identifies a <i>HOX</i> -Centric Network Associated with Serous Ovarian Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1574-1584.	1.1	28
141	Spatiotemporal estimation of historical PM 2.5 concentrations using PM 10, meteorological variables, and spatial effect. Atmospheric Environment, 2017, 166, 182-191.	1.9	28
142	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. BMC Medicine, 2020, 18, 229.	2.3	28
143	Reproductive factors, hormone use and gastric cancer risk: The Singapore Chinese Health Study. International Journal of Cancer, 2016, 138, 2837-2845.	2.3	27
144	Systematic meta-analyses, field synopsis and global assessment of the evidence of genetic association studies in colorectal cancer. Gut, 2020, 69, 1460-1471.	6.1	27

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145	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. American Journal of Clinical Nutrition, 2021, 113, 1490-1502.	2.2	27
146	Polygenic risk scores for prediction of breast cancer risk in Asian populations. Genetics in Medicine, 2022, 24, 586-600.	1.1	27
147	Enhanced <i>GAB2</i> Expression Is Associated with Improved Survival in High-Grade Serous Ovarian Cancer and Sensitivity to PI3K Inhibition. Molecular Cancer Therapeutics, 2015, 14, 1495-1503.	1.9	26
148	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 860-870.	1.1	26
149	Population-based targeted sequencing of 54 candidate genes identifies <i>PALB2</i> as a susceptibility gene for high-grade serous ovarian cancer. Journal of Medical Genetics, 2021, 58, 305-313.	1.5	26
150	Birth weight and other prenatal factors and risk of breast cancer in Asian-Americans. Breast Cancer Research and Treatment, 2011, 130, 917-925.	1.1	25
151	Cigarette smoking is associated with adverse survival among women with ovarian cancer: Results from a pooled analysis of 19 studies. International Journal of Cancer, 2017, 140, 2422-2435.	2.3	25
152	Stomach Cancer Disparity among Korean Americans by Tumor Characteristics: Comparison with Non-Hispanic Whites, Japanese Americans, South Koreans, and Japanese. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 587-596.	1.1	25
153	Identification of Novel Loci and New Risk Variant in Known Loci for Colorectal Cancer Risk in East Asians. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 477-486.	1.1	25
154	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). Journal of Genetics and Genome Research, 2015, 2, .	0.3	25
155	Genetic Variation in <i>TYMS</i> in the One-Carbon Transfer Pathway Is Associated with Ovarian Carcinoma Types in the Ovarian Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 1822-1830.	1.1	24
156	Family history of cancer and gastroesophageal disorders and risk of esophageal and gastric adenocarcinomas: a case–control study. BMC Cancer, 2014, 14, 60.	1,1	24
157	A Newly Identified Susceptibility Locus near <i>FOXP1</i> Modifies the Association of Gastroesophageal Reflux with Barrett's Esophagus. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1739-1747.	1.1	24
158	Fine-Scale Mapping of the 4q24 Locus Identifies Two Independent Loci Associated with Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1680-1691.	1.1	24
159	Green tea intake is associated with urinary estrogen profiles in Japanese-American women. Nutrition Journal, 2013, 12, 25.	1.5	23
160	Enrichment of putative PAX8 target genes at serous epithelial ovarian cancer susceptibility loci. British Journal of Cancer, 2017, 116, 524-535.	2.9	23
161	A pooled analysis of dietary sugar/carbohydrate intake and esophageal and gastric cardia adenocarcinoma incidence and survival in the USA. International Journal of Epidemiology, 2017, 46, 1836-1846.	0.9	23
162	Associations Between Reproductive and Hormone-Related Factors and Risk of Nonalcoholic Fatty Liver Disease in a Multiethnic Population. Clinical Gastroenterology and Hepatology, 2021, 19, 1258-1266.e1.	2.4	23

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163	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362.	1.4	23
164	Soy intake and other lifestyle determinants of serum estrogen levels among postmenopausal Chinese women in Singapore. Cancer Epidemiology Biomarkers and Prevention, 2002, 11, 844-51.	1.1	23
165	Genetic variation in insulin-like growth factor 2 may play a role in ovarian cancer risk. Human Molecular Genetics, 2011, 20, 2263-2272.	1.4	22
166	Epithelialâ€Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. Genetic Epidemiology, 2015, 39, 689-697.	0.6	22
167	The Effect of Reduced Dietary Fat and Soy Supplementation on Circulating Adipocytokines in Postmenopausal Women: A Randomized Controlled 2-Month Trial. Nutrition and Cancer, 2016, 68, 554-559.	0.9	22
168	Timing of births and oral contraceptive use influences ovarian cancer risk. International Journal of Cancer, 2017, 141, 2392-2399.	2.3	22
169	Association between genetically predicted polycystic ovary syndrome and ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2019, 48, 822-830.	0.9	22
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