

Hoda Anton-Culver

List of Publications by Year in descending order

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Version: 2024-02-01

229
papers

18,375
citations

18436

62
h-index

16127

124
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232
all docs

232
docs citations

232
times ranked

23204
citing authors

#	ARTICLE	IF	CITATIONS
1	Antibodies to Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) in <i>All of Us</i> Research Program Participants, 2 January to 18 March 2020. <i>Clinical Infectious Diseases</i> , 2022, 74, 584-590.	2.9	26
2	High Prediagnosis Inflammation-Related Risk Score Associated with Decreased Ovarian Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 443-452.	1.1	2
3	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
4	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
5	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
6	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women. <i>Scientific Reports</i> , 2022, 12, 6199.	1.6	2
7	Relationship between BMI trajectories and cardiometabolic outcomes in postmenopausal women: a growth mixture modeling approach. <i>Annals of Epidemiology</i> , 2022, 72, 9-17.	0.9	2
8	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
9	Expanding Our Understanding of Ovarian Cancer Risk: The Role of Incomplete Pregnancies. <i>Journal of the National Cancer Institute</i> , 2021, 113, 301-308.	3.0	8
10	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	2.9	5
11	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , 2021, 384, 440-451.	13.9	414
12	Differences in Melanoma Between Canada and New South Wales, Australia: A Population-Based Genes, Environment, and Melanoma (GEM) Study. <i>JID Innovations</i> , 2021, 1, 100002.	1.2	1
13	Metabolically Healthy/Unhealthy Overweight/Obesity Associations With Incident Heart Failure in Postmenopausal Women. <i>Circulation: Heart Failure</i> , 2021, 14, e007297.	1.6	7
14	Comparison of Perioperative Outcomes for Radical Nephrectomy Based on Surgical Approach for Masses Greater than 10cm. <i>Journal of Endourology</i> , 2021, 35, 1785-1792.	1.1	2
15	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203.	2.6	6
16	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021, 23, 86.	2.2	7
17	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145.	2.9	9
18	The WISDOM study: a new approach to screening can and should be tested. <i>Breast Cancer Research and Treatment</i> , 2021, 189, 593-598.	1.1	19

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19	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	1.1	19
20	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021, 11, 19787.	1.6	2
21	Germline Pathogenic Variants in Cancer Predisposition Genes Among Women With Invasive Lobular Carcinoma of the Breast. <i>Journal of Clinical Oncology</i> , 2021, 39, 3918-3926.	0.8	22
22	Disease-Associated Risk Variants in <i>ANRIL</i> Are Associated with Tumor-Infiltrating Lymphocyte Presence in Primary Melanomas in the Population-Based GEM Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 2309-2316.	1.1	2
23	Endometriosis and menopausal hormone therapy impact the hysterectomy-ovarian cancer association. <i>Gynecologic Oncology</i> , 2021, , .	0.6	5
24	Association of Melanoma-Risk Variants with Primary Melanoma Tumor Prognostic Characteristics and Melanoma-Specific Survival in the GEM Study. <i>Current Oncology</i> , 2021, 28, 4756-4771.	0.9	1
25	Mammography screening and mortality by risk status in the California teachers study. <i>BMC Cancer</i> , 2021, 21, 1341.	1.1	4
26	Inherited Melanoma Risk Variants Associated with Histopathologically Amelanotic Melanoma. <i>Journal of Investigative Dermatology</i> , 2020, 140, 918-922.e7.	0.3	1
27	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
28	A California Cancer Registry Analysis of Urothelial and Non-urothelial Bladder Cancer Subtypes: Epidemiology, Treatment, and Survival. <i>Clinical Genitourinary Cancer</i> , 2020, 18, e330-e336.	0.9	12
29	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	2.6	39
30	Association of Known Melanoma Risk Factors with Primary Melanoma of the Scalp and Neck. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2203-2210.	1.1	6
31	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581.	9.4	265
32	Contribution of Germline Predisposition Gene Mutations to Breast Cancer Risk in African American Women. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1213-1221.	3.0	51
33	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	1.6	2
34	Sociodemographic and metabolic risk characteristics associated with metabolic weight categories in the Women's Health Initiative. <i>Cardiovascular Endocrinology and Metabolism</i> , 2020, 9, 42-48.	0.5	2
35	Incidence of diabetes according to metabolically healthy or unhealthy normal weight or overweight/obesity in postmenopausal women: the Women's Health Initiative. <i>Menopause</i> , 2020, 27, 640-647.	0.8	16
36	Menopausal hormone therapy prior to the diagnosis of ovarian cancer is associated with improved survival. <i>Gynecologic Oncology</i> , 2020, 158, 702-709.	0.6	15

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37	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
38	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
39	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
40	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	2.3	28
41	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
42	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
43	Secondhand smoke, obesity, and risk of type II diabetes among California teachers. <i>Annals of Epidemiology</i> , 2019, 32, 35-42.	0.9	9
44	Association between genetically predicted polycystic ovary syndrome and ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2019, 48, 822-830.	0.9	22
45	Racial and Socioeconomic Disparities in Bladder Cancer Survival: Analysis of the California Cancer Registry. <i>Clinical Genitourinary Cancer</i> , 2019, 17, e995-e1002.	0.9	34
46	Evaluation of vitamin D biosynthesis and pathway target genes reveals UGT2A1/2 and EGFR polymorphisms associated with epithelial ovarian cancer in African American Women. <i>Cancer Medicine</i> , 2019, 8, 2503-2513.	1.3	6
47	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
48	Associations of sleep duration with cardiometabolic outcomes in American Indians and Alaska Natives and other race/ethnicities: results from the BRFSS. <i>Sleep Health</i> , 2019, 5, 344-351.	1.3	19
49	MC1R variants in childhood and adolescent melanoma: a retrospective pooled analysis of a multicentre cohort. <i>The Lancet Child and Adolescent Health</i> , 2019, 3, 332-342.	2.7	16
50	A breast cancer case-control study of polybrominated diphenyl ether (PBDE) serum levels among California women. <i>Environment International</i> , 2019, 127, 412-419.	4.8	28
51	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	2.9	52
52	Relationship of Chromosome Arm 10q Variants to Occurrence of Multiple Primary Melanoma in the Population-Based Genes, Environment, and Melanoma (GEM) Study. <i>Journal of Investigative Dermatology</i> , 2019, 139, 1410-1412.	0.3	0
53	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	2.6	711
54	Genetic susceptibility to radiation-induced breast cancer after Hodgkin lymphoma. <i>Blood</i> , 2019, 133, 1130-1139.	0.6	29

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55	A comprehensive gene-environment interaction analysis in Ovarian Cancer using genome-wide significant common variants. <i>International Journal of Cancer</i> , 2019, 144, 2192-2205.	2.3	12
56	The <i>BRCA2</i> c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018, 39, 729-741.	1.1	19
57	Identification of a gene expression signature predicting survival in oral cavity squamous cell carcinoma using Monte Carlo cross validation. <i>Oral Oncology</i> , 2018, 78, 72-79.	0.8	2
58	Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. <i>British Journal of Cancer</i> , 2018, 118, 1123-1129.	2.9	15
59	Cancer surveillance in northern Africa, and central and western Asia: challenges and strategies in support of developing cancer registries. <i>Lancet Oncology</i> , The, 2018, 19, e85-e92.	5.1	34
60	36-Item Short Form Survey (SF-36) Versus Gait Speed As Predictor of Preclinical Mobility Disability in Older Women: The Women's Health Initiative. <i>Journal of the American Geriatrics Society</i> , 2018, 66, 706-713.	1.3	13
61	Perception matters: Stressful life events increase breast cancer risk. <i>Journal of Psychosomatic Research</i> , 2018, 110, 46-53.	1.2	14
62	Robust Tests for Additive Gene-Environment Interaction in Case-Control Studies Using Gene-Environment Independence. <i>American Journal of Epidemiology</i> , 2018, 187, 366-377.	1.6	8
63	The interaction between vitamin D receptor polymorphisms and sun exposure around time of diagnosis influences melanoma survival. <i>Pigment Cell and Melanoma Research</i> , 2018, 31, 287-296.	1.5	13
64	Time Trends in Per- and Polyfluoroalkyl Substances (PFASs) in California Women: Declining Serum Levels, 2011-2015. <i>Environmental Science & Technology</i> , 2018, 52, 277-287.	4.6	54
65	Negative Valence Life Events Promote Breast Cancer Development. <i>Clinical Breast Cancer</i> , 2018, 18, e521-e528.	1.1	5
66	Polycystic Ovary Syndrome, Oligomenorrhea, and Risk of Ovarian Cancer Histotypes: Evidence from the Ovarian Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 174-182.	1.1	20
67	Trends in Treatment Patterns and Clinical Outcomes in Young Women Diagnosed With Ductal Carcinoma In Situ. <i>Clinical Breast Cancer</i> , 2018, 18, e179-e185.	1.1	17
68	Breast Cancer Characteristics in Middle Eastern Women Immigrants Compared With Non-Hispanic White Women in California. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky014.	1.4	5
69	Breast cancer risk and serum levels of per- and poly-fluoroalkyl substances: a case-control study nested in the California Teachers Study. <i>Environmental Health</i> , 2018, 17, 83.	1.7	48
70	Diet Quality Scores Inversely Associated with Postmenopausal Breast Cancer Risk Are Not Associated with Premenopausal Breast Cancer Risk in the California Teachers Study. <i>Journal of Nutrition</i> , 2018, 148, 1830-1837.	1.3	24
71	Inherited Genetic Variants Associated with Melanoma BRAF/NRAS Subtypes. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2398-2404.	0.3	9
72	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2018, 78, 5419-5430.	0.4	54

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73	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. PLoS ONE, 2018, 13, e0197561.	1.1	9
74	A Case-Control Study of the Genetic Variability in Reactive Oxygen Speciesâ€™Metabolizing Enzymes in Melanoma Risk. International Journal of Molecular Sciences, 2018, 19, 242.	1.8	10
75	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. International Journal of Molecular Sciences, 2018, 19, 2473.	1.8	3
76	Laminin 332 expression and prognosis in breast cancer. Human Pathology, 2018, 82, 289-296.	1.1	18
77	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. Nature Genetics, 2018, 50, 968-978.	9.4	184
78	Enrichment of putative PAX8 target genes at serous epithelial ovarian cancer susceptibility loci. British Journal of Cancer, 2017, 116, 524-535.	2.9	23
79	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
80	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799.	0.4	75
81	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	9.4	356
82	Use of common analgesic medications and ovarian cancer survival: results from a pooled analysis in the Ovarian Cancer Association Consortium. British Journal of Cancer, 2017, 116, 1223-1228.	2.9	13
83	Temporal Evaluation of Polybrominated Diphenyl Ether (PBDE) Serum Levels in Middle-Aged and Older California Women, 2011â€“2015. Environmental Science & Technology, 2017, 51, 4697-4704.	4.6	55
84	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
85	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
86	Genome-Wide Testing of Exonic Variants and Breast Cancer Risk in the California Teachers Study. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1462-1465.	1.1	0
87	Associations of MC1R Genotype and Patient Phenotypes with BRAF and NRAS Mutations in Melanoma. Journal of Investigative Dermatology, 2017, 137, 2588-2598.	0.3	11
88	History of Comorbidities and Survival of Ovarian Cancer Patients, Results from the Ovarian Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1470-1473.	1.1	10
89	Association of Incident Amelanotic Melanoma With Phenotypic Characteristics, <i>MC1R</i> Status, and Prior Amelanotic Melanoma. JAMA Dermatology, 2017, 153, 1026.	2.0	19
90	Cancer risk in different generations of <i>Middle Eastern</i> immigrants to <i>California</i> , 1988â€“2013. International Journal of Cancer, 2017, 141, 2260-2269.	2.3	7

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91	Sex differences in the association of cutaneous melanoma incidence rates and geographic ultraviolet light exposure. <i>Journal of the American Academy of Dermatology</i> , 2017, 76, 499-505.e3.	0.6	66
92	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	1.1	67
93	No Evidence That Genetic Variation in the Myeloid-Derived Suppressor Cell Pathway Influences Ovarian Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 420-424.	1.1	3
94	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	0.8	9
95	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. <i>Oncotarget</i> , 2016, 7, 80140-80163.	0.8	31
96	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	3.9	118
97	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	1.1	12
98	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 884-895.	0.9	71
99	Fine-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
100	Exome genotyping arrays to identify rare and low frequency variants associated with epithelial ovarian cancer risk. <i>Human Molecular Genetics</i> , 2016, 25, 3600-3612.	1.4	17
101	Nevus count associations with pigmented phenotype, histopathological melanoma characteristics and survival from melanoma. <i>International Journal of Cancer</i> , 2016, 139, 1217-1222.	2.3	11
102	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	1.5	174
103	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	2.2	31
104	Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. <i>Human Genetics</i> , 2016, 135, 741-756.	1.8	19
105	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016, 27, 679-693.	0.8	21
106	Cancer burden in four countries of the Middle East Cancer Consortium (Cyprus; Jordan; Israel; Izmir) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 <i>Cancer Epidemiology</i> , 2016, 44, 195-202.	0.8	16
107	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 1619-1630.	0.9	111
108	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

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109	Sex Differences in the Relationship between Fitness and Obesity on Risk for Asthma in Adolescents. <i>Journal of Pediatrics</i> , 2016, 176, 36-42.	0.9	35
110	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
111	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
112	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
113	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). <i>Scientific Reports</i> , 2016, 6, 32512.	1.6	19
114	Variants in autophagy-related genes and clinical characteristics in melanoma: a population-based study. <i>Cancer Medicine</i> , 2016, 5, 3336-3345.	1.3	23
115	Novel polymorphisms in caspase-8 are associated with breast cancer risk in the California Teachers Study. <i>BMC Cancer</i> , 2016, 16, 14.	1.1	18
116	Treatment for T1a Renal Cancer Substratified by Size: "Less is More". <i>Journal of Urology</i> , 2016, 196, 1000-1007.	0.2	26
117	Preliminary Associations between the Detection of Perfluoroalkyl Acids (PFAAs) in Drinking Water and Serum Concentrations in a Sample of California Women. <i>Environmental Science and Technology Letters</i> , 2016, 3, 264-269.	3.9	50
118	Assessment of Multifactor Gene-Environment Interactions and Ovarian Cancer Risk: Candidate Genes, Obesity, and Hormone-Related Risk Factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 780-790.	1.1	10
119	Association of Interferon Regulatory Factor-4 Polymorphism rs12203592 With Divergent Melanoma Pathways. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw004.	3.0	28
120	The association between socioeconomic status and tumour stage at diagnosis of ovarian cancer: A pooled analysis of 18 case-control studies. <i>Cancer Epidemiology</i> , 2016, 41, 71-79.	0.8	20
121	No evidence that protein truncating variants in <i>BRIP1</i> are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016, 53, 298-309.	1.5	94
122	Breast cancer risk variants at 6q25 display different phenotype associations and regulate <i>ESR1</i> , <i>RMND1</i> and <i>CCDC170</i> . <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
123	Investigation of Exomic Variants Associated with Overall Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 446-454.	1.1	9
124	Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. <i>Human Genetics</i> , 2016, 135, 137-154.	1.8	8
125	Vitamin D receptor polymorphisms and survival in patients with cutaneous melanoma: a population-based study. <i>Carcinogenesis</i> , 2016, 37, 30-38.	1.3	54
126	No clinical utility of <i>KRAS</i> variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18

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127	Duration of Adulthood Overweight, Obesity, and Cancer Risk in the Women's Health Initiative: A Longitudinal Study from the United States. <i>PLoS Medicine</i> , 2016, 13, e1002081.	3.9	99
128	Supplementing Public Health Inspection via Social Media. <i>PLoS ONE</i> , 2016, 11, e0152117.	1.1	29
129	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.	1.1	26
130	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. <i>Oncotarget</i> , 2016, 7, 69097-69110.	0.8	5
131	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. <i>Oncotarget</i> , 2016, 7, 72381-72394.	0.8	13
132	A targeted genetic association study of epithelial ovarian cancer susceptibility. <i>Oncotarget</i> , 2016, 7, 7381-7389.	0.8	7
133	Epithelial-Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. <i>Genetic Epidemiology</i> , 2015, 39, 689-697.	0.6	22
134	Clinical Implementation of a Breast Cancer Risk Assessment Program in a Multiethnic Patient Population: Which Risk Model to Use?. <i>Breast Journal</i> , 2015, 21, 562-564.	0.4	6
135	Inherited Variation at MC1R and Histological Characteristics of Primary Melanoma. <i>PLoS ONE</i> , 2015, 10, e0119920.	1.1	22
136	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. <i>PLoS ONE</i> , 2015, 10, e0128106.	1.1	44
137	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	428
138	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	1.4	40
139	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
140	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. <i>Carcinogenesis</i> , 2015, 36, 256-271.	1.3	14
141	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
142	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
143	Genome-wide significant risk associations for mucinous ovarian carcinoma. <i>Nature Genetics</i> , 2015, 47, 888-897.	9.4	78
144	Network-Based Integration of GWAS and Gene Expression Identifies a HOX-Centric Network Associated with Serous Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1574-1584.	1.1	28

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145	Socioeconomic status as a predictor of adherence to treatment guidelines for early-stage ovarian cancer. <i>Gynecologic Oncology</i> , 2015, 138, 121-127.	0.6	49
146	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. <i>Clinical Cancer Research</i> , 2015, 21, 5264-5276.	3.2	33
147	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , 2015, 97, 22-34.	2.6	37
148	Impact of race, socioeconomic status, and the health care system on the treatment of advanced-stage ovarian cancer in California. <i>American Journal of Obstetrics and Gynecology</i> , 2015, 212, 468.e1-468.e9.	0.7	73
149	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. <i>Gynecologic Oncology</i> , 2015, 136, 542-548.	0.6	15
150	Impact of National Cancer Institute Comprehensive Cancer Centers on Ovarian Cancer Treatment and Survival. <i>Journal of the American College of Surgeons</i> , 2015, 220, 940-950.	0.2	94
151	Inherited Genetic Variants Associated with Occurrence of Multiple Primary Melanoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 992-997.	1.1	36
152	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. <i>Nature Communications</i> , 2015, 6, 8234.	5.8	63
153	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.	9.4	357
154	Common variants at the <i>CHEK2</i> gene locus and risk of epithelial ovarian cancer. <i>Carcinogenesis</i> , 2015, 36, 1341-1353.	1.3	24
155	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. <i>Human Molecular Genetics</i> , 2015, 24, 5955-5964.	1.4	68
156	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv219.	3.0	99
157	Fine-Scale Mapping of the 4q24 Locus Identifies Two Independent Loci Associated with Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1680-1691.	1.1	24
158	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, 285-298.	1.4	38
159	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). <i>Journal of Genetics and Genome Research</i> , 2015, 2, .	0.3	25
160	Human Nail Clippings as a Source of DNA for Genetic Studies. <i>Open Journal of Epidemiology</i> , 2015, 05, 41-50.	0.2	8
161	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	1.1	49
162	Sun Exposure and Melanoma Survival: A GEM Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2145-2152.	1.1	26

#	ARTICLE	IF	CITATIONS
163	Comparison of Clinicopathologic Features and Survival of Histopathologically Amelanotic and Pigmented Melanomas. <i>JAMA Dermatology</i> , 2014, 150, 1306.	2.0	142
164	Association between Class III Obesity (BMI of 40â€“59 kg/m ²) and Mortality: A Pooled Analysis of 20 Prospective Studies. <i>PLoS Medicine</i> , 2014, 11, e1001673.	3.9	299
165	Spatial analysis of adherence to treatment guidelines for advanced-stage ovarian cancer and the impact of race and socioeconomic status. <i>Gynecologic Oncology</i> , 2014, 134, 60-67.	0.6	99
166	High-volume ovarian cancer care: Survival impact and disparities in access for advanced-stage disease. <i>Gynecologic Oncology</i> , 2014, 132, 403-410.	0.6	141
167	Attitudes Toward Cancer Clinical Trial Participation in Young Adults with a History of Cancer and a Healthy College Student Sample: A Preliminary Investigation. <i>Journal of Adolescent and Young Adult Oncology</i> , 2014, 3, 20-27.	0.7	25
168	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. <i>Journal of the National Cancer Institute</i> , 2014, 106, djt431-djt431.	3.0	186
169	Alcohol Consumption and Survival after a Breast Cancer Diagnosis: A Literature-Based Meta-analysis and Collaborative Analysis of Data for 29,239 Cases. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 934-945.	1.1	37
170	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014, 514, 92-97.	13.7	548
171	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
172	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014, 23, 6034-6046.	1.4	12
173	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. <i>Breast Cancer Research</i> , 2014, 16, R51.	2.2	14
174	Self-recalled Youth Physical Activity and Postmenopausal Cardiovascular Disease. <i>Health Behavior and Policy Review</i> , 2014, 1, 472-483.	0.3	1
175	Multilevel socioeconomic effects on quality of life in adolescent and young adult survivors of leukemia and lymphoma. <i>Quality of Life Research</i> , 2013, 22, 1339-1351.	1.5	19
176	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	326
177	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
178	Tumor-Infiltrating Lymphocyte Grade in Primary Melanomas Is Independently Associated With Melanoma-Specific Survival in the Population-Based Genes, Environment and Melanoma Study. <i>Journal of Clinical Oncology</i> , 2013, 31, 4252-4259.	0.8	232
179	A unique gender difference in early onset melanoma implies that in addition to ultraviolet light exposure other causative factors are important. <i>Pigment Cell and Melanoma Research</i> , 2013, 26, 128-135.	1.5	48
180	Genome-wide association studies identify four ER negativeâ€“specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374

#	ARTICLE	IF	CITATIONS
181	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
182	A genome-wide association study to identify genetic susceptibility loci that modify ductal and lobular postmenopausal breast cancer risk associated with menopausal hormone therapy use: a two-stage design with replication. <i>Breast Cancer Research and Treatment</i> , 2013, 138, 529-542.	1.1	18
183	Evidence of Gene-Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. <i>PLoS Genetics</i> , 2013, 9, e1003284.	1.5	136
184	Combined and Interactive Effects of Environmental and GWAS-Identified Risk Factors in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 880-890.	1.1	54
185	Obesity and risk of ovarian cancer subtypes: evidence from the Ovarian Cancer Association Consortium. <i>Endocrine-Related Cancer</i> , 2013, 20, 251-262.	1.6	169
186	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , 2013, 4, 1628.	5.8	144
187	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , 2013, 4, 1627.	5.8	98
188	Relation between self-recalled childhood physical activity and adult physical activity: The women's health initiative. <i>Open Journal of Epidemiology</i> , 2013, 03, 224-231.	0.2	2
189	<i>CHEK2</i> 1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancer-Specific Death, and Increased Risk of a Second Breast Cancer. <i>Journal of Clinical Oncology</i> , 2012, 30, 4308-4316.	0.8	162
190	Meat consumption, ornithine decarboxylase gene polymorphism, and outcomes after colorectal cancer diagnosis. <i>Journal of Carcinogenesis</i> , 2012, 11, 17.	2.5	6
191	Association between endometriosis and risk of histological subtypes of ovarian cancer: a pooled analysis of case-control studies. <i>Lancet Oncology</i> , The, 2012, 13, 385-394.	5.1	753
192	Association between insurance and socioeconomic status and risk of advanced stage Hodgkin lymphoma in adolescents and young adults. <i>Cancer</i> , 2012, 118, 6179-6187.	2.0	74
193	Vitamin D receptor polymorphisms in patients with cutaneous melanoma. <i>International Journal of Cancer</i> , 2012, 130, 405-418.	2.3	61
194	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 3289-3303.	1.4	152
195	The association of polymorphisms in hormone metabolism pathway genes, menopausal hormone therapy, and breast cancer risk: a nested case-control study in the California Teachers Study cohort. <i>Breast Cancer Research</i> , 2011, 13, R37.	2.2	15
196	Clinically Relevant Changes in Family History of Cancer Over Time. <i>JAMA - Journal of the American Medical Association</i> , 2011, 306, 172-8.	3.8	40
197	The association of body mass index with mortality in the California Teachers Study. <i>International Journal of Cancer</i> , 2011, 129, 2492-2501.	2.3	33
198	Common variants at 19p13 are associated with susceptibility to ovarian cancer. <i>Nature Genetics</i> , 2010, 42, 880-884.	9.4	235

#	ARTICLE	IF	CITATIONS
199	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. <i>Nature Genetics</i> , 2010, 42, 874-879.	9.4	321
200	Socioeconomic Impacts on Survival Differ by Race/Ethnicity among Adolescents and Young Adults with Non-Hodgkin's Lymphoma. <i>Journal of Cancer Epidemiology</i> , 2010, 2010, 1-10.	0.5	39
201	Genetic Variation in <i>TYMS</i> in the One-Carbon Transfer Pathway Is Associated with Ovarian Carcinoma Types in the Ovarian Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1822-1830.	1.1	24
202	Associations of Cumulative Sun Exposure and Phenotypic Characteristics with Histologic Solar Elastosis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2932-2941.	1.1	45
203	Kids, Adolescents, and Young Adults Cancer Study—A Methodologic Approach in Cancer Epidemiology Research. <i>Journal of Cancer Epidemiology</i> , 2009, 2009, 1-7.	0.5	4
204	Associations of a Polymorphism in the Ornithine Decarboxylase Gene with Colorectal Cancer Survival. <i>Clinical Cancer Research</i> , 2009, 15, 6208-6216.	3.2	35
205	Leukemia survival in children, adolescents, and young adults: influence of socioeconomic status and other demographic factors. <i>Cancer Causes and Control</i> , 2009, 20, 1409-1420.	0.8	52
206	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. <i>Nature Genetics</i> , 2009, 41, 996-1000.	9.4	276
207	Accurate classification of <i>MLH1/MSH2</i> missense variants with multivariate analysis of protein polymorphisms-mismatch repair (MAPP-MMR). <i>Human Mutation</i> , 2008, 29, 852-860.	1.1	101
208	Survival After Colorectal Cancer Diagnosis Is Associated with Colorectal Cancer Family History. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3134-3140.	1.1	32
209	Testing Targeted Approaches to Enhance Cancer Genetics Network Minority Recruitment within Asian Populations. <i>Public Health Genomics</i> , 2008, 11, 234-240.	1.0	12
210	Differential Effects of Wine Consumption on Colorectal Cancer Outcomes Based on Family History of the Disease. <i>Nutrition and Cancer</i> , 2007, 59, 36-45.	0.9	28
211	Risk and risk reduction involving arginine intake and meat consumption in colorectal tumorigenesis and survival. <i>International Journal of Cancer</i> , 2007, 120, 459-468.	2.3	68
212	CDKN2A Germline Mutations in Individuals with Cutaneous Malignant Melanoma. <i>Journal of Investigative Dermatology</i> , 2007, 127, 1234-1243.	0.3	50
213	The Prevalence of CDKN2A Germ-Line Mutations and Relative Risk for Cutaneous Malignant Melanoma: An International Population-Based Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1520-1525.	1.1	105
214	Low Allele Frequency of MLH1 D132H in American Colorectal and Endometrial Cancer Patients. <i>Diseases of the Colon and Rectum</i> , 2005, 48, 1723-1727.	0.7	14
215	Lifetime Risk of Melanoma in CDKN2A Mutation Carriers in a Population-Based Sample. <i>Journal of the National Cancer Institute</i> , 2005, 97, 1507-1515.	3.0	200
216	Aurora-A/STK15 T + 91A is a general low penetrance cancer susceptibility gene: a meta-analysis of multiple cancer types. <i>Carcinogenesis</i> , 2005, 26, 1368-1373.	1.3	132

#	ARTICLE	IF	CITATIONS
217	Rrp1b, a new candidate susceptibility gene for breast cancer progression and metastasis. PLoS Genetics, 2005, preprint, e214.	1.5	0
218	Validation of family history data in cancer family registries. American Journal of Preventive Medicine, 2003, 24, 190-198.	1.6	220
219	The Cancer Genetics Network: Recruitment Results and Pilot Studies. Public Health Genomics, 2003, 6, 171-177.	0.6	51
220	Characteristics and survival of patients with non-Hodgkin's lymphoma with and without acquired immunodeficiency syndrome. Hematological Oncology, 2002, 20, 177-187.	0.8	8
221	Recent diet and breast cancer risk: the California Teachers Study (USA). Cancer Causes and Control, 2002, 13, 407-415.	0.8	185
222	High breast cancer incidence rates among California teachers: results from the California Teachers Study (United States). Cancer Causes and Control, 2002, 13, 625-635.	0.8	206
223	RESPONSE: Re: Characterization of Hereditary Nonpolyposis Colorectal Cancer Families From a Population-Based Series of Cases. Journal of the National Cancer Institute, 2001, 93, 717-717.	3.0	0
224	Let's Leave the Date Out of the Name of the Standard Population. Journal of the National Cancer Institute, 2000, 92, 1269-1270.	3.0	2
225	Characterization of Hereditary Nonpolyposis Colorectal Cancer Families From a Population-Based Series of Cases. Journal of the National Cancer Institute, 2000, 92, 1517-1522.	3.0	80
226	Differences in breast cancer stage at diagnosis between non-Hispanic white and Hispanic populations, San Diego County 1988-1993. Breast Cancer Research and Treatment, 1998, 50, 1-9.	1.1	49
227	Unique features of gastric carcinoma in the young. , 1998, 83, 25-33.		67
228	Validation of family history of breast cancer and identification of the BRCA1 and other syndromes using a population-based cancer registry. Genetic Epidemiology, 1996, 13, 193-205.	0.6	45
229	Sex differences in lung-cancer risk associated with cigarette smoking. International Journal of Cancer, 1993, 54, 44-48.	2.3	153