

# Esther M John

## List of Publications by Year in descending order

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Version: 2024-02-01

250  
papers

21,819  
citations

16411

64  
h-index

11581

135  
g-index

261  
all docs

261  
docs citations

261  
times ranked

22019  
citing authors

#	ARTICLE	IF	CITATIONS
1	Maternal and prenatal factors and age at thelarche in the LEGACY Girls Study cohort: implications for breast cancer risk. <i>International Journal of Epidemiology</i> , 2023, 52, 272-283.	0.9	1
2	Smoking, Radiation Therapy, and Contralateral Breast Cancer Risk in Young Women. <i>Journal of the National Cancer Institute</i> , 2022, 114, 631-634.	3.0	6
3	Performance of African-ancestry-specific polygenic hazard score varies according to local ancestry in 8q24. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, 25, 229-237.	2.0	9
4	Breast and Prostate Cancer Risks for Male <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variant Carriers Using Polygenic Risk Scores. <i>Journal of the National Cancer Institute</i> , 2022, 114, 109-122.	3.0	19
5	Cumulative menstrual months and breast cancer risk by hormone receptor status and ethnicity: The Breast Cancer Etiology in Minorities Study. <i>International Journal of Cancer</i> , 2022, 150, 208-220.	2.3	0
6	Risks of breast and ovarian cancer for women harboring pathogenic missense variants in <i>BRCA1</i> and <i>BRCA2</i> compared with those harboring protein truncating variants. <i>Genetics in Medicine</i> , 2022, 24, 119-129.	1.1	10
7	Predictors of urinary polycyclic aromatic hydrocarbon metabolites in girls from the San Francisco Bay Area. <i>Environmental Research</i> , 2022, 205, 112534.	3.7	4
8	Cancer Risks Associated With <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>Journal of Clinical Oncology</i> , 2022, 40, 1529-1541.	0.8	90
9	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
10	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
11	A Rare Germline <i>HOXB13</i> Variant Contributes to Risk of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , 2022, 81, 458-462.	0.9	22
12	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
13	Oral Contraceptive Use in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Absolute Cancer Risks and Benefits. <i>Journal of the National Cancer Institute</i> , 2022, 114, 540-552.	3.0	7
14	OUP accepted manuscript. <i>International Journal of Epidemiology</i> , 2022, . .	0.9	0
15	Improvement on recovery and reproducibility for quantifying urinary mono-hydroxylated polycyclic aromatic hydrocarbons (OH-PAHs). <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2022, 1192, 123113.	1.2	4
16	Prostate cancer risk stratification improvement across multiple ancestries with new polygenic hazard score. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, 25, 755-761.	2.0	14
17	Differences in Thickness-Specific Incidence and Factors Associated With Cutaneous Melanoma in the US From 2010 to 2018. <i>JAMA Oncology</i> , 2022, 8, 755.	3.4	20
18	Polygenic risk scores for prediction of breast cancer risk in Asian populations. <i>Genetics in Medicine</i> , 2022, 24, 586-600.	1.1	27

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19	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traitsâ€”The Hispanic/Latino Anthropometry Consortium. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100099.	1.0	3
20	Association of contralateral breast cancer risk with mammographic density defined at higherâ€”thanâ€”conventional intensity thresholds. <i>International Journal of Cancer</i> , 2022, 151, 1304-1309.	2.3	3
21	Weight is More Informative than Body Mass Index for Predicting Postmenopausal Breast Cancer Risk: Prospective Family Study Cohort (ProF-SC). <i>Cancer Prevention Research</i> , 2022, 15, 185-191.	0.7	4
22	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. <i>Breast Cancer Research</i> , 2022, 24, 27.	2.2	15
23	Polygenic risk scores for prediction of breast cancer risk in women of African ancestry: a cross-ancestry approach. <i>Human Molecular Genetics</i> , 2022, 31, 3133-3143.	1.4	11
24	Relevance of the MHC region for breast cancer susceptibility in Asians. <i>Breast Cancer</i> , 2022, 29, 869-879.	1.3	1
25	Overall survival is the lowest among young women with postpartum breast cancer. <i>European Journal of Cancer</i> , 2022, 168, 119-127.	1.3	10
26	Breast cancer diagnosis and treatment during the COVID-19 pandemic in a nationwide, insured population. <i>Breast Cancer Research and Treatment</i> , 2022, 194, 475-482.	1.1	14
27	Adherence to the 2020 American Cancer Society Guideline for Cancer Prevention and risk of breast cancer for women at increased familial and genetic risk in the Breast Cancer Family Registry: an evaluation of the weight, physical activity, and alcohol consumption recommendations. <i>Breast Cancer Research and Treatment</i> , 2022, 194, 673-682.	1.1	1
28	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
29	Comparing 5-Year and Lifetime Risks of Breast Cancer Using the Prospective Family Study Cohort. <i>Journal of the National Cancer Institute</i> , 2021, 113, 785-791.	3.0	13
30	Africanâ€”specific improvement of a polygenic hazard score for age at diagnosis of prostate cancer. <i>International Journal of Cancer</i> , 2021, 148, 99-105.	2.3	24
31	The Impact of the first COVID-19 shelter-in-place announcement on social distancing, difficulty in daily activities, and levels of concern in the San Francisco Bay Area: A cross-sectional social media survey. <i>PLoS ONE</i> , 2021, 16, e0244819.	1.1	5
32	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
33	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , 2021, 53, 65-75.	9.4	264
34	Additional SNPs improve risk stratification of a polygenic hazard score for prostate cancer. <i>Prostate Cancer and Prostatic Diseases</i> , 2021, 24, 532-541.	2.0	16
35	Race, ethnicity and risk of second primary contralateral breast cancer in the United States. <i>International Journal of Cancer</i> , 2021, 148, 2748-2758.	2.3	13
36	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	5.8	19

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37	Polygenic hazard score is associated with prostate cancer in multi-ethnic populations. <i>Nature Communications</i> , 2021, 12, 1236.	5.8	40
38	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
39	Association of Risk-Reducing Salpingo-Oophorectomy With Breast Cancer Risk in Women With BRCA1 and BRCA2 Pathogenic Variants. <i>JAMA Oncology</i> , 2021, 7, 585-592.	3.4	16
40	Evaluating Polygenic Risk Scores for Breast Cancer in Women of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2021, 113, 1168-1176.	3.0	41
41	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	2.6	18
42	Treatment and Monitoring Variability in US Metastatic Breast Cancer Care. <i>JCO Clinical Cancer Informatics</i> , 2021, 5, 600-614.	1.0	5
43	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous BRCA1 or BRCA2 pathogenic variant. <i>Genetics in Medicine</i> , 2021, 23, 1726-1737.	1.1	16
44	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. <i>Nature Communications</i> , 2021, 12, 4198.	5.8	24
45	A competing risks model with binary time varying covariates for estimation of breast cancer risks in BRCA1 families. <i>Statistical Methods in Medical Research</i> , 2021, 30, 2165-2183.	0.7	2
46	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
47	Performance of the IBIS/Tyler-Cuzick model of breast cancer risk by race and ethnicity in the Women's Health Initiative. <i>Cancer</i> , 2021, 127, 3742-3750.	2.0	21
48	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
49	Risk of Breast Cancer Among Carriers of Pathogenic Variants in Breast Cancer Predisposition Genes Varies by Polygenic Risk Score. <i>Journal of Clinical Oncology</i> , 2021, 39, 2564-2573.	0.8	47
50	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
51	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
52	Coronary Artery Disease in Young Women After Radiation Therapy for Breast Cancer. <i>JACC: CardioOncology</i> , 2021, 3, 381-392.	1.7	31
53	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
54	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

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55	Racial/Ethnic Disparities in Survival after Breast Cancer Diagnosis by Estrogen and Progesterone Receptor Status: A Pooled Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 351-363.	1.1	7
56	Mammographic texture features associated with contralateral breast cancer in the WECARE Study. <i>Npj Breast Cancer</i> , 2021, 7, 146.	2.3	1
57	Recreational Physical Activity and Outcomes After Breast Cancer in Women at High Familial Risk. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab090.	1.4	1
58	Association of Genomic Domains in <i>BRCA1</i> and <i>BRCA2</i> with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , 2020, 80, 624-638.	0.4	39
59	A genome-wide association study of prostate cancer in Latinos. <i>International Journal of Cancer</i> , 2020, 146, 1819-1826.	2.3	24
60	Considerations When Using Breast Cancer Risk Models for Women with Negative <i>BRCA1/BRCA2</i> Mutation Results. <i>Journal of the National Cancer Institute</i> , 2020, 112, 418-422.	3.0	1
61	Recreational Physical Activity Is Associated with Reduced Breast Cancer Risk in Adult Women at High Risk for Breast Cancer: A Cohort Study of Women Selected for Familial and Genetic Risk. <i>Cancer Research</i> , 2020, 80, 116-125.	0.4	37
62	A Polygenic Risk Score for Breast Cancer in US Latinas and Latin American Women. <i>Journal of the National Cancer Institute</i> , 2020, 112, 590-598.	3.0	53
63	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
64	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	1.1	82
65	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. <i>Nature Communications</i> , 2020, 11, 3905.	5.8	28
66	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833.	5.8	88
67	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	2.6	39
68	Association of germline variation with the survival of women with <i>BRCA1/2</i> pathogenic variants and breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 44.	2.3	5
69	A case-control study of the joint effect of reproductive factors and radiation treatment for first breast cancer and risk of contralateral breast cancer in the WECARE study. <i>Breast</i> , 2020, 54, 62-69.	0.9	3
70	The <i>CHEK2</i> Variant C.349A>G Is Associated with Prostate Cancer Risk and Carriers Share a Common Ancestor. <i>Cancers</i> , 2020, 12, 3254.	1.7	16
71	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
72	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

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73	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , 2020, 78, 316-320.	0.9	32
74	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
75	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.	5.8	46
76	Characterization of the Cancer Spectrum in Men With Germline <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>JAMA Oncology</i> , 2020, 6, 1218.	3.4	48
77	Radiation Treatment, <i>ATM</i> , <i>BRCA1/2</i> , and <i>CHEK2</i> *1100delC Pathogenic Variants and Risk of Contralateral Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1275-1279.	3.0	21
78	Menstrual and reproductive characteristics and breast cancer risk by hormone receptor status and ethnicity: The Breast Cancer Etiology in Minorities study. <i>International Journal of Cancer</i> , 2020, 147, 1808-1822.	2.3	10
79	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
80	Alcohol Consumption, Cigarette Smoking, and Risk of Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Results from The <i>BRCA1</i> and <i>BRCA2</i> Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 368-378.	1.1	24
81	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
82	Risk-reducing salpingo-oophorectomy, natural menopause, and breast cancer risk: an international prospective cohort of <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research</i> , 2020, 22, 8.	2.2	41
83	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. <i>Blood Advances</i> , 2020, 4, 181-190.	2.5	16
84	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 781-794.	0.9	37
85	Surveillance of cancer among sexual and gender minority populations: Where are we and where do we need to go?. <i>Cancer</i> , 2019, 125, 4360-4362.	2.0	10
86	Re-evaluating genetic variants identified in candidate gene studies of breast cancer risk using data from nearly 280,000 women of Asian and European ancestry. <i>EBioMedicine</i> , 2019, 48, 203-211.	2.7	14
87	Two truncating variants in <i>FANCC</i> and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
88	Estrogenic activity, race/ethnicity, and Indigenous American ancestry among San Francisco Bay Area women. <i>PLoS ONE</i> , 2019, 14, e0213809.	1.1	4
89	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. <i>JAMA Network Open</i> , 2019, 2, e1912259.	2.8	5
90	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88

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91	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	2.9	19
92	The functional ALDH2 polymorphism is associated with breast cancer risk: A pooled analysis from the Breast Cancer Association Consortium. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2019, 7, e707.	0.6	9
93	Regular use of aspirin and other non-steroidal anti-inflammatory drugs and breast cancer risk for women at familial or genetic risk: a cohort study. <i>Breast Cancer Research</i> , 2019, 21, 52.	2.2	44
94	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
95	Enrollment and biospecimen collection in a multiethnic family cohort: the Northern California site of the Breast Cancer Family Registry. <i>Cancer Causes and Control</i> , 2019, 30, 395-408.	0.8	13
96	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
97	Association of Prepubertal and Adolescent Androgen Concentrations With Timing of Breast Development and Family History of Breast Cancer. <i>JAMA Network Open</i> , 2019, 2, e190083.	2.8	7
98	Benign breast disease increases breast cancer risk independent of underlying familial risk profile: Findings from a Prospective Family Study Cohort. <i>International Journal of Cancer</i> , 2019, 145, 370-379.	2.3	9
99	10-year performance of four models of breast cancer risk: a validation study. <i>Lancet Oncology</i> , The, 2019, 20, 504-517.	5.1	116
100	Race/Ethnicity and Accuracy of Self-Reported Female First-Degree Family History of Breast and Other Cancers in the Northern California Breast Cancer Family Registry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1792-1801.	1.1	4
101	Quantitative Ultrasound Spectroscopy for Differentiation of Hepatocellular Carcinoma from At-Risk and Normal Liver Parenchyma. <i>Clinical Cancer Research</i> , 2019, 25, 6683-6691.	3.2	8
102	Alcohol consumption, cigarette smoking, and familial breast cancer risk: findings from the Prospective Family Study Cohort (ProF-SC). <i>Breast Cancer Research</i> , 2019, 21, 128.	2.2	27
103	A Pooled Analysis of Breastfeeding and Breast Cancer Risk by Hormone Receptor Status in Parous Hispanic Women. <i>Epidemiology</i> , 2019, 30, 449-457.	1.2	10
104	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
105	Identification of novel common breast cancer risk variants at the 6q25 locus among Latinas. <i>Breast Cancer Research</i> , 2019, 21, 3.	2.2	32
106	Risk-Reducing Oophorectomy and Breast Cancer Risk Across the Spectrum of Familial Risk. <i>Journal of the National Cancer Institute</i> , 2019, 111, 331-334.	3.0	31
107	Obesity, Body Composition, and Breast Cancer. <i>JAMA Oncology</i> , 2018, 4, 804.	3.4	14
108	Intake of bean fiber, beans, and grains and reduced risk of hormone receptor-negative breast cancer: the San Francisco Bay Area Breast Cancer Study. <i>Cancer Medicine</i> , 2018, 7, 2131-2144.	1.3	23



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109	Mutational spectrum in a worldwide study of 29,700 families with <i>BRCA1</i> or <i>BRCA2</i> mutations. <i>Human Mutation</i> , 2018, 39, 593-620.	1.1	224
110	Reproductive history, breastfeeding and risk of triple negative breast cancer: The Breast Cancer Etiology in Minorities (BEM) study. <i>International Journal of Cancer</i> , 2018, 142, 2273-2285.	2.3	56
111	Impact of individual and neighborhood factors on disparities in prostate cancer survival. <i>Cancer Epidemiology</i> , 2018, 53, 1-11.	0.8	61
112	Discovery of mutations in homologous recombination genes in African-American women with breast cancer. <i>Familial Cancer</i> , 2018, 17, 187-195.	0.9	3
113	Breast Cancer Family History and Contralateral Breast Cancer Risk in Young Women: An Update From the Women's Environmental Cancer and Radiation Epidemiology Study. <i>Journal of Clinical Oncology</i> , 2018, 36, 1513-1520.	0.8	44
114	The Influence of Number and Timing of Pregnancies on Breast Cancer Risk for Women With <i>BRCA1</i> or <i>BRCA2</i> Mutations. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky078.	1.4	21
115	Age-specific breast cancer risk by body mass index and familial risk: prospective family study cohort (ProF-SC). <i>Breast Cancer Research</i> , 2018, 20, 132.	2.2	51
116	CYP2D6 phenotype, tamoxifen, and risk of contralateral breast cancer in the WECARE Study. <i>Breast Cancer Research</i> , 2018, 20, 149.	2.2	11
117	Oral Contraceptive Use and Breast Cancer Risk: Retrospective and Prospective Analyses From a <i>BRCA1</i> and <i>BRCA2</i> Mutation Carrier Cohort Study. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky023.	1.4	33
118	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , 2018, 13, e0196245.	1.1	9
119	Metabolomic profiles in breast cancer: a pilot case-control study in the breast cancer family registry. <i>BMC Cancer</i> , 2018, 18, 532.	1.1	17
120	The association of mammographic density with risk of contralateral breast cancer and change in density with treatment in the WECARE study. <i>Breast Cancer Research</i> , 2018, 20, 23.	2.2	24
121	Impact of individual and neighborhood factors on socioeconomic disparities in localized and advanced prostate cancer risk. <i>Cancer Causes and Control</i> , 2018, 29, 951-966.	0.8	24
122	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018, 50, 928-936.	9.4	652
123	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018, 9, 2256.	5.8	88
124	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	9.4	184
125	Germline Variation and Breast Cancer Incidence: A Gene-Based Association Study and Whole-Genome Prediction of Early-Onset Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 1057-1064.	1.1	9
126	Response to Conner et al. Re: "Cigarette Smoking and Breast Cancer Risk in Hispanic and Non-Hispanic White Women: The Breast Cancer Health Disparities Study". <i>Journal of Women's Health</i> , 2017, 26, 92-93.	1.5	1



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127	Limited influence of germline genetic variation on all-cause mortality in women with early onset breast cancer: evidence from gene-based tests, single-marker regression, and whole-genome prediction. <i>Breast Cancer Research and Treatment</i> , 2017, 164, 707-717.	1.1	4
128	Association of Common Genetic Variants With Contralateral Breast Cancer Risk in the WECARE Study. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	3.0	28
129	Alcohol consumption and cigarette smoking in combination: A predictor of contralateral breast cancer risk in the WECARE study. <i>International Journal of Cancer</i> , 2017, 141, 916-924.	2.3	31
130	Risks of Breast, Ovarian, and Contralateral Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>JAMA - Journal of the American Medical Association</i> , 2017, 317, 2402.	3.8	1,898
131	Reply to Dietary isoflavone intake and all-cause mortality in breast cancer survivors: The <i>Breast Cancer Family Registry</i> methodological issues. <i>Cancer</i> , 2017, 123, 3639-3639.	2.0	1
132	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
133	The Interaction between Genetic Ancestry and Breast Cancer Risk Factors among Hispanic Women: The Breast Cancer Health Disparities Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 692-701.	1.1	19
134	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
135	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
136	Panel sequencing of 264 candidate susceptibility genes and segregation analysis in a cohort of non- <i>BRCA1</i> , non- <i>BRCA2</i> breast cancer families. <i>Breast Cancer Research and Treatment</i> , 2017, 166, 937-949.	1.1	16
137	Association of breast cancer risk in <i>BRCA1</i> and <i>BRCA2</i> mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 117-134.	1.1	18
138	Pre-diagnostic breastfeeding, adiposity, and mortality among parous Hispanic and non-Hispanic white women with invasive breast cancer: the Breast Cancer Health Disparities Study. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 321-331.	1.1	4
139	Genetic modifiers of <i>CHEK2</i> *1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	1.1	67
140	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	3.0	57
141	Hormone receptor status of a first primary breast cancer predicts contralateral breast cancer risk in the WECARE study population. <i>Breast Cancer Research</i> , 2017, 19, 83.	2.2	27
142	Assessing biological and technological variability in protein levels measured in pre-diagnostic plasma samples of women with breast cancer. <i>Biomarker Research</i> , 2017, 5, 30.	2.8	13
143	Prediction of Breast and Prostate Cancer Risks in Male <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers Using Polygenic Risk Scores. <i>Journal of Clinical Oncology</i> , 2017, 35, 2240-2250.	0.8	152
144	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

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