

Per Hall

List of Publications by Year in descending order

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Version: 2024-02-01

256
papers

27,843
citations

18482

62
h-index

6996

154
g-index

267
all docs

267
docs citations

267
times ranked

36445
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	27.8	3,823
2	Risk of Ischemic Heart Disease in Women after Radiotherapy for Breast Cancer. <i>New England Journal of Medicine</i> , 2013, 368, 987-998.	27.0	3,028
3	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	21.4	2,634
4	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	21.4	1,818
5	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
6	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	21.4	960
7	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014, 514, 92-97.	27.8	548
8	Breast Cancer Risk Genes – Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	27.0	532
9	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.	21.4	513
10	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.	21.4	493
11	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	428
12	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017, 49, 834-841.	21.4	426
13	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	21.4	374
14	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.	21.4	357
15	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015, 11, e1005378.	3.5	331
16	Genomic Diversity and Admixture Differs for Stone-Age Scandinavian Foragers and Farmers. <i>Science</i> , 2014, 344, 747-750.	12.6	315
17	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
18	Effect of low doses of ionising radiation in infancy on cognitive function in adulthood: Swedish population based cohort study. <i>BMJ: British Medical Journal</i> , 2004, 328, 19-0.	2.3	280

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19	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.	21.4	265
20	Treatment-Associated Leukemia Following Testicular Cancer. <i>Journal of the National Cancer Institute</i> , 2000, 92, 1165-1171.	6.3	225
21	Intrinsic molecular signature of breast cancer in a population-based cohort of 412 patients. <i>Breast Cancer Research</i> , 2006, 8, R34.	5.0	218
22	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	6.2	201
23	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	21.4	184
24	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	27.8	183
25	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	12.8	178
26	Personalized early detection and prevention of breast cancer: ENVISION consensus statement. <i>Nature Reviews Clinical Oncology</i> , 2020, 17, 687-705.	27.6	178
27	<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.	3.2	174
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. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	9.4	157
29	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.	2.9	152
30	Evidence of Gene-Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. <i>PLoS Genetics</i> , 2013, 9, e1003284.	3.5	136
31	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.	21.4	125
32	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
33	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.	8.4	118
34	Mammographic Density Reduction Is a Prognostic Marker of Response to Adjuvant Tamoxifen Therapy in Postmenopausal Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2013, 31, 2249-2256.	1.6	113
35	Thyroid cancer risk after thyroid examination with 131I: A population-based cohort study in Sweden. <i>International Journal of Cancer</i> , 2003, 106, 580-587.	5.1	112
36	Common variants in <i>ZNF365</i> are associated with both mammographic density and breast cancer risk. <i>Nature Genetics</i> , 2011, 43, 185-187.	21.4	109

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37	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. <i>Nature Communications</i> , 2014, 5, 5303.	12.8	109
38	Cancer mortality after iodine-131 therapy for hyperthyroidism. <i>International Journal of Cancer</i> , 1992, 50, 886-890.	5.1	107
39	Combined genetic and splicing analysis of BRCA1 c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. <i>Human Molecular Genetics</i> , 2016, 25, 2256-2268.	2.9	106
40	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	12.8	105
41	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166.	2.5	101
42	Risk Factors and Tumor Characteristics of Interval Cancers by Mammographic Density. <i>Journal of Clinical Oncology</i> , 2015, 33, 1030-1037.	1.6	99
43	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv219.	6.3	99
44	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.	6.2	98
45	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.	3.2	94
46	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	12.8	93
47	A combined analysis of genome-wide association studies in breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011, 126, 717-727.	2.5	90
48	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	12.8	90
49	Cohort Profile: The Karolinska Mammography Project for Risk Prediction of Breast Cancer (KARMA). <i>International Journal of Epidemiology</i> , 2017, 46, 1740-1741g.	1.9	88
50	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. <i>International Journal of Epidemiology</i> , 2018, 47, 526-536.	1.9	88
51	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
52	Cardiac doses from Swedish breast cancer radiotherapy since the 1950s. <i>Radiotherapy and Oncology</i> , 2009, 90, 127-135.	0.6	87
53	Assessment of Breast Cancer Risk Factors Reveals Subtype Heterogeneity. <i>Cancer Research</i> , 2017, 77, 3708-3717.	0.9	87
54	Association of ESR1 gene tagging SNPs with breast cancer risk. <i>Human Molecular Genetics</i> , 2009, 18, 1131-1139.	2.9	84

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55	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. <i>Breast Cancer Research</i> , 2010, 12, R110.	5.0	82
56	A clinical model for identifying the short-term risk of breast cancer. <i>Breast Cancer Research</i> , 2017, 19, 29.	5.0	79
57	Predictors of Discontinuation of Adjuvant Hormone Therapy in Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2015, 33, 2262-2269.	1.6	78
58	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	12.8	78
59	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016, 48, 667-674.	21.4	77
60	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.	6.2	76
61	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.9	75
62	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020, 11, 3353.	12.8	75
63	Incidence and survival of Swedish patients with differentiated thyroid cancer. <i>International Journal of Cancer</i> , 2003, 106, 569-573.	5.1	71
64	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	2.4	67
65	Non-Hodgkin's lymphoma and skin malignancies-shared etiology?. <i>International Journal of Cancer</i> , 1995, 62, 519-522.	5.1	64
66	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1503-1510.	2.5	64
67	Automated Measurement of Volumetric Mammographic Density: A Tool for Widespread Breast Cancer Risk Assessment. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1764-1772.	2.5	62
68	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.	3.1	62
69	Genetic overlap between endometriosis and endometrial cancer: evidence from cross-disease genetic correlation and GWAS meta-analyses. <i>Cancer Medicine</i> , 2018, 7, 1978-1987.	2.8	62
70	Differences in mammographic density between Asian and Caucasian populations: a comparative analysis. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 353-362.	2.5	61
71	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.	6.2	59
72	Independent risk of depression, anxiety, and stress-related disorders in patients with invasive and <i>in situ</i> breast cancer. <i>International Journal of Cancer</i> , 2017, 140, 841-852.	5.1	59

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73	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	56
74	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , 2015, 75, 2457-2467.	0.9	55
75	Mammographic Breast Density and Breast Cancer: Evidence of a Shared Genetic Basis. <i>Cancer Research</i> , 2012, 72, 1478-1484.	0.9	54
76	Prediction of breast cancer risk based on common genetic variants in women of East Asian ancestry. <i>Breast Cancer Research</i> , 2016, 18, 124.	5.0	52
77	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	6.4	52
78	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.	5.1	51
79	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744.	7.1	51
80	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 1478-1492.	2.9	50
81	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	2.5	49
82	Hormone-replacement therapy influences gene expression profiles and is associated with breast-cancer prognosis: a cohort study. <i>BMC Medicine</i> , 2006, 4, 16.	5.5	47
83	Increased Risk of Second Primary Malignancies in Patients with Gynecological Cancer: A Swedish record-linkage study. <i>Acta Oncologica</i> , 1995, 34, 771-777.	1.8	45
84	Associations of Breast Cancer Risk Prediction Tools With Tumor Characteristics and Metastasis. <i>Journal of Clinical Oncology</i> , 2016, 34, 251-258.	1.6	45
85	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	1.9	45
86	A comprehensive tool for measuring mammographic density changes over time. <i>Breast Cancer Research and Treatment</i> , 2018, 169, 371-379.	2.5	45
87	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.	6.3	45
88	Change of mammographic density predicts the risk of contralateral breast cancer - a case-control study. <i>Breast Cancer Research</i> , 2013, 15, R57.	5.0	44
89	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016, 18, 22.	5.0	43
90	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017, 19, 119.	5.0	43

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91	Common diseases as determinants of menopausal age. <i>Human Reproduction</i> , 2016, 31, 2856-2864.	0.9	42
92	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	2.9	40
93	Identification of Women at High Risk of Breast Cancer Who Need Supplemental Screening. <i>Radiology</i> , 2020, 297, 327-333.	7.3	40
94	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.	3.5	39
95	Patient survival and tumor characteristics associated with CHEK2:p.1157T → findings from the Breast Cancer Association Consortium. <i>Breast Cancer Research</i> , 2016, 18, 98.	5.0	39
96	Public interest in and acceptability of the prospect of risk-stratified screening for breast and prostate cancer. <i>Acta Oncologica</i> , 2016, 55, 45-51.	1.8	39
97	Women's perceptions of personalized risk-based breast cancer screening and prevention: An international focus group study. <i>Psycho-Oncology</i> , 2019, 28, 1056-1062.	2.3	39
98	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	6.2	39
99	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.	2.9	38
100	Influence of Lifestyle Factors on Mammographic Density in Postmenopausal Women. <i>PLoS ONE</i> , 2013, 8, e81876.	2.5	37
101	Risk of esophageal cancer following radiotherapy for Hodgkin lymphoma. <i>Haematologica</i> , 2014, 99, e193-e196.	3.5	37
102	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.	6.2	37
103	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369.	3.3	35
104	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. <i>International Journal of Cancer</i> , 2021, 148, 307-319.	5.1	35
105	Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. <i>International Journal of Cancer</i> , 2015, 136, E685-96.	5.1	34
106	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. <i>Human Molecular Genetics</i> , 2016, 25, 3863-3876.	2.9	33
107	Low-Dose Tamoxifen for Mammographic Density Reduction: A Randomized Controlled Trial. <i>Journal of Clinical Oncology</i> , 2021, 39, 1899-1908.	1.6	33
108	Implantable catheter system for long-term intravenous chemotherapy. <i>Journal of Surgical Oncology</i> , 1989, 41, 39-41.	1.7	32

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109	Mammographic Density Change and Risk of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 391-399.	6.3	32
110	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	1.3	32
111	Mammographic microcalcifications and risk of breast cancer. <i>British Journal of Cancer</i> , 2021, 125, 759-765.	6.4	32
112	Radiation-Associated Thyroid Cancer - Facts and Fiction. <i>Acta Oncologica</i> , 1998, 37, 325-330.	1.8	31
113	Proteomics profiling identify CAPS as a potential predictive marker of tamoxifen resistance in estrogen receptor positive breast cancer. <i>Clinical Proteomics</i> , 2015, 12, 8.	2.1	31
114	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.	1.8	31
115	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.	5.0	31
116	Time-dependent risk and predictors of venous thromboembolism in breast cancer patients: A population-based cohort study. <i>Cancer</i> , 2017, 123, 468-475.	4.1	31
117	Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. <i>Breast Cancer Research</i> , 2019, 21, 68.	5.0	31
118	Prevalence of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants in a large, unselected breast cancer cohort. <i>International Journal of Cancer</i> , 2019, 144, 1195-1204.	5.1	31
119	CYP2D6 Genotype Predicts Tamoxifen Discontinuation and Prognosis in Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2020, 38, 548-557.	1.6	31
120	Increased pancreatic cancer risk following radiotherapy for testicular cancer. <i>British Journal of Cancer</i> , 2016, 115, 901-908.	6.4	30
121	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	12.8	30
122	Tumors after Radiotherapy for Thyroid Cancer; A case-control study within a cohort of thyroid cancer patients. <i>Acta Oncologica</i> , 1992, 31, 403-407.	1.8	28
123	Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Gene-Environment Interactions. <i>Genetic Epidemiology</i> , 2014, 38, 84-93.	1.3	28
124	Determinants of Mammographic Density Change. <i>JNCI Cancer Spectrum</i> , 2019, 3, pkz004.	2.9	27
125	Genetic modifiers of menopausal hormone replacement therapy and breast cancer risk: a genome-wide interaction study. <i>Endocrine-Related Cancer</i> , 2013, 20, 875-887.	3.1	26
126	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	5.0	26

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127	RAD51B in Familial Breast Cancer. PLoS ONE, 2016, 11, e0153788.	2.5	26
128	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. Endocrine-Related Cancer, 2015, 22, 851-861.	3.1	25
129	Interval breast cancer is associated with other types of tumors. Nature Communications, 2019, 10, 4648.	12.8	25
130	Evaluation of Exome Sequencing to Estimate Tumor Burden in Plasma. PLoS ONE, 2014, 9, e104417.	2.5	25
131	Area and Volumetric Density Estimation in Processed Full-Field Digital Mammograms for Risk Assessment of Breast Cancer. PLoS ONE, 2014, 9, e110690.	2.5	24
132	Identification of two novel mammographic density loci at 6Q25.1. Breast Cancer Research, 2015, 17, 75.	5.0	24
133	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.	2.5	24
134	Association of Microcalcification Clusters with Short-term Invasive Breast Cancer Risk and Breast Cancer Risk Factors. Scientific Reports, 2019, 9, 14604.	3.3	24
135	Prediction and clinical utility of a contralateral breast cancer risk model. Breast Cancer Research, 2019, 21, 144.	5.0	24
136	Worse quality of life in young and recently diagnosed breast cancer survivors compared with female survivors of other cancers: A cross-sectional study. International Journal of Cancer, 2016, 139, 2415-2425.	5.1	23
137	Disease trajectories and mortality among women diagnosed with breast cancer. Breast Cancer Research, 2019, 21, 95.	5.0	23
138	Infection-related hospitalizations in breast cancer patients: Risk and impact on prognosis. Journal of Infection, 2016, 72, 650-658.	3.3	22
139	Risk of hospitalisation and death due to bone fractures after breast cancer: a registry-based cohort study. British Journal of Cancer, 2016, 115, 1400-1407.	6.4	22
140	Comparison of handheld ultrasound and automated breast ultrasound in women recalled after mammography screening. Acta Radiologica, 2017, 58, 515-520.	1.1	22
141	Aspirin and other non-steroidal anti-inflammatory drugs and depression, anxiety, and stress-related disorders following a cancer diagnosis: a nationwide register-based cohort study. BMC Medicine, 2020, 18, 238.	5.5	22
142	Heritability of Mammographic Breast Density, Density Change, Microcalcifications, and Masses. Cancer Research, 2020, 80, 1590-1600.	0.9	22
143	Volumetric Mammographic Density: Heritability and Association With Breast Cancer Susceptibility Loci. Journal of the National Cancer Institute, 2014, 106, dju334-dju334.	6.3	21
144	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. Cancer Causes and Control, 2016, 27, 679-693.	1.8	21

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145	The HLA-DQ ² 1 insertion is a strong achalasia risk factor and displays a geospatial north-south gradient among Europeans. <i>European Journal of Human Genetics</i> , 2016, 24, 1228-1231.	2.8	21
146	SNP-SNP interaction analysis of NF- κ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , 2015, 6, 37979-37994.	1.8	20
147	Gene-environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. <i>International Journal of Cancer</i> , 2017, 141, 1830-1840.	5.1	20
148	Hormonal determinants of mammographic density and density change. <i>Breast Cancer Research</i> , 2020, 22, 95.	5.0	20
149	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. <i>Breast Cancer Research</i> , 2015, 17, 110.	5.0	19
150	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.	3.3	19
151	The <i>BRCA2</i> c.68-7T variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018, 39, 729-741.	2.5	19
152	Differential Burden of Rare and Common Variants on Tumor Characteristics, Survival, and Mode of Detection in Breast Cancer. <i>Cancer Research</i> , 2018, 78, 6329-6338.	0.9	19
153	Mammographic density change in a cohort of premenopausal women receiving tamoxifen for breast cancer prevention over 5 years. <i>Breast Cancer Research</i> , 2020, 22, 101.	5.0	19
154	European women's perceptions of the implementation and organisation of risk-based breast cancer screening and prevention: a qualitative study. <i>BMC Cancer</i> , 2020, 20, 247.	2.6	19
155	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	12.8	19
156	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.	2.5	19
157	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51.	8.2	19
158	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.	2.5	18
159	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	1.4	18
160	Common genetic variation and novel loci associated with volumetric mammographic density. <i>Breast Cancer Research</i> , 2018, 20, 30.	5.0	18
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