

Celine M Vachon

List of Publications by Year in descending order

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Version: 2024-02-01

279
papers

20,392
citations

12330

69
h-index

12946

131
g-index

283
all docs

283
docs citations

283
times ranked

19510
citing authors

#	ARTICLE	IF	CITATIONS
1	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	21.4	960
3	Benign Breast Disease and the Risk of Breast Cancer. <i>New England Journal of Medicine</i> , 2005, 353, 229-237.	27.0	785
4	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	6.2	711
5	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. <i>Journal of the National Cancer Institute</i> , 2011, 103, 250-263.	6.3	596
6	Inherited Mutations in 17 Breast Cancer Susceptibility Genes Among a Large Triple-Negative Breast Cancer Cohort Unselected for Family History of Breast Cancer. <i>Journal of Clinical Oncology</i> , 2015, 33, 304-311.	1.6	521
7	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
8	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
9	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
10	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , 2021, 384, 440-451.	27.0	414
11	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	21.4	374
12	Unsupervised Deep Learning Applied to Breast Density Segmentation and Mammographic Risk Scoring. <i>IEEE Transactions on Medical Imaging</i> , 2016, 35, 1322-1331.	8.9	360
13	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	3.5	315
14	Association of mammographically defined percent breast density with epidemiologic risk factors for breast cancer (United States). <i>Cancer Causes and Control</i> , 2000, 11, 653-662.	1.8	311
15	A locus on 19p13 modifies risk of breast cancer in BRCA1 mutation carriers and is associated with hormone receptor-negative breast cancer in the general population. <i>Nature Genetics</i> , 2010, 42, 885-892.	21.4	309
16	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
17	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. <i>Nature Genetics</i> , 2011, 43, 1210-1214.	21.4	279
18	Mammographic density, breast cancer risk and risk prediction. <i>Breast Cancer Research</i> , 2007, 9, 217.	5.0	270

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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	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. <i>Journal of the National Cancer Institute</i> , 2014, 106, .	6.3	261
21	Prevention of Breast Cancer in Postmenopausal Women: Approaches to Estimating and Reducing Risk. <i>Journal of the National Cancer Institute</i> , 2009, 101, 384-398.	6.3	226
22	Age-Related Lobular Involution and Risk of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2006, 98, 1600-1607.	6.3	218
23	Mammographic Breast Density and Subsequent Risk of Breast Cancer in Postmenopausal Women According to Tumor Characteristics. <i>Journal of the National Cancer Institute</i> , 2011, 103, 1179-1189.	6.3	192
24	Breast Cancer Risk by Breast Density, Menopause, and Postmenopausal Hormone Therapy Use. <i>Journal of Clinical Oncology</i> , 2010, 28, 3830-3837.	1.6	188
25	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
26	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	27.8	183
27	Mammographic Breast Density as a General Marker of Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 43-49.	2.5	181
28	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876.	21.4	179
29	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	12.8	178
30	The Contributions of Breast Density and Common Genetic Variation to Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	174
31	<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
32	Breast Density and Benign Breast Disease: Risk Assessment to Identify Women at High Risk of Breast Cancer. <i>Journal of Clinical Oncology</i> , 2015, 33, 3137-3143.	1.6	170
33	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012, 21, 5373-5384.	2.9	168
34	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
35	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for <i>CH</i> <i>EK</i> <i>2</i> *110delC Carriers. <i>Journal of Clinical Oncology</i> , 2016, 34, 2750-2760.	1.6	152
36	Mammographic density and risk of breast cancer by age and tumor characteristics. <i>Breast Cancer Research</i> , 2013, 15, R104.	5.0	146

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37	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. <i>Carcinogenesis</i> , 2014, 35, 1012-1019.	2.8	145
38	Comparison of Clinical and Automated Breast Density Measurements: Implications for Risk Prediction and Supplemental Screening. <i>Radiology</i> , 2016, 279, 710-719.	7.3	145
39	Breast cancer risk prediction using a clinical risk model and polygenic risk score. <i>Breast Cancer Research and Treatment</i> , 2016, 159, 513-525.	2.5	129
40	Increased risk of monoclonal gammopathy in first-degree relatives of patients with multiple myeloma or monoclonal gammopathy of undetermined significance. <i>Blood</i> , 2009, 114, 785-790.	1.4	127
41	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	21.4	125
42	Texture Features from Mammographic Images and Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 837-845.	2.5	121
43	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
44	Longitudinal Trends in Mammographic Percent Density and Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 921-928.	2.5	118
45	Genome-wide association study identifies a novel susceptibility locus at 6p21.3 among familial CLL. <i>Blood</i> , 2011, 117, 1911-1916.	1.4	118
46	A new statistic for identifying batch effects in high-throughput genomic data that uses guided principal component analysis. <i>Bioinformatics</i> , 2013, 29, 2877-2883.	4.1	118
47	Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. <i>Nature Genetics</i> , 2011, 43, 185-187.	21.4	109
48	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
49	Common Genetic Variation and Breast Cancer Risk—Past, Present, and Future. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 380-394.	2.5	108
50	Mammographic density and ageing: A collaborative pooled analysis of cross-sectional data from 22 countries worldwide. <i>PLoS Medicine</i> , 2017, 14, e1002335.	8.4	108
51	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	12.8	105
52	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
53	19p13.1 Is a Triple-Negative-Specific Breast Cancer Susceptibility Locus. <i>Cancer Research</i> , 2012, 72, 1795-1803.	0.9	100
54	Benign Breast Disease, Mammographic Breast Density, and the Risk of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2013, 105, 1043-1049.	6.3	99

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55	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
56	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
57	Awareness of Breast Density and Its Impact on Breast Cancer Detection and Risk. <i>Journal of Clinical Oncology</i> , 2015, 33, 1143-1150.	1.6	96
58	Assessment of the Accuracy of the Gail Model in Women With Atypical Hyperplasia. <i>Journal of Clinical Oncology</i> , 2008, 26, 5374-5379.	1.6	94
59	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	12.8	94
60	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
61	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	12.8	93
62	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
63	Age-specific Trends in Mammographic Density: The Minnesota Breast Cancer Family Study. <i>American Journal of Epidemiology</i> , 2008, 167, 1027-1036.	3.4	88
64	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
65	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
66	Incidence of AL Amyloidosis in Olmsted County, Minnesota, 1990 through 2015. <i>Mayo Clinic Proceedings</i> , 2019, 94, 465-471.	3.0	87
67	Association Between Mammographic Density and Age-Related Lobular Involution of the Breast. <i>Journal of Clinical Oncology</i> , 2010, 28, 2207-2212.	1.6	84
68	Breast Density and Breast Cancer Risk: A Practical Review. <i>Mayo Clinic Proceedings</i> , 2014, 89, 548-557.	3.0	84
69	Breast cancer risk in women with radial scars in benign breast biopsies. <i>Breast Cancer Research and Treatment</i> , 2008, 108, 167-174.	2.5	83
70	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
71	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
72	Tissue composition of mammographically dense and non-dense breast tissue. <i>Breast Cancer Research and Treatment</i> , 2012, 131, 267-275.	2.5	72

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73	Strong Evidence of a Genetic Determinant for Mammographic Density, a Major Risk Factor for Breast Cancer. <i>Cancer Research</i> , 2007, 67, 8412-8418.	0.9	69
74	Familial monoclonal gammopathy of undetermined significance and multiple myeloma: epidemiology, risk factors, and biological characteristics. <i>Blood</i> , 2012, 119, 5359-5366.	1.4	68
75	Association of Genetic Variation in Genes Implicated in the β -Catenin Destruction Complex with Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2101-2108.	2.5	67
76	An Automated Approach for Estimation of Breast Density. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3090-3097.	2.5	67
77	A Novel Automated Mammographic Density Measure and Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2012, 104, 1028-1037.	6.3	67
78	Independent Association of Lobular Involution and Mammographic Breast Density With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2010, 102, 1716-1723.	6.3	66
79	Radiomic Phenotypes of Mammographic Parenchymal Complexity: Toward Augmenting Breast Density in Breast Cancer Risk Assessment. <i>Radiology</i> , 2019, 290, 41-49.	7.3	63
80	Common occurrence of monoclonal B-cell lymphocytosis among members of high-risk CLL families. <i>British Journal of Haematology</i> , 2010, 151, 152-158.	2.5	61
81	Complex fibroadenoma and breast cancer risk: a Mayo Clinic Benign Breast Disease Cohort Study. <i>Breast Cancer Research and Treatment</i> , 2015, 153, 397-405.	2.5	61
82	Detection and prevalence of monoclonal gammopathy of undetermined significance: a study utilizing mass spectrometry-based monoclonal immunoglobulin rapid accurate mass measurement. <i>Blood Cancer Journal</i> , 2019, 9, 102.	6.2	57
83	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	56
84	Germline Lysine-Specific Demethylase 1 (<i>LSD1/KDM1A</i>) Mutations Confer Susceptibility to Multiple Myeloma. <i>Cancer Research</i> , 2018, 78, 2747-2759.	0.9	56
85	Automated and Clinical Breast Imaging Reporting and Data System Density Measures Predict Risk for Screen-Detected and Interval Cancers. <i>Annals of Internal Medicine</i> , 2018, 168, 757-765.	3.9	56
86	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
87	Mammographic Breast Density and Breast Cancer: Evidence of a Shared Genetic Basis. <i>Cancer Research</i> , 2012, 72, 1478-1484.	0.9	54
88	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	2.9	53
89	Association of Mammographic Density with the Pathology of Subsequent Breast Cancer among Postmenopausal Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 872-879.	2.5	52
90	Body mass index, mammographic density, and breast cancer risk by estrogen receptor subtype. <i>Breast Cancer Research</i> , 2019, 21, 48.	5.0	52

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91	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
92	Model for Individualized Prediction of Breast Cancer Risk After a Benign Breast Biopsy. <i>Journal of Clinical Oncology</i> , 2015, 33, 923-929.	1.6	51
93	Annexin A1 expression in a pooled breast cancer series: association with tumor subtypes and prognosis. <i>BMC Medicine</i> , 2015, 13, 156.	5.5	51
94	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	2.5	49
95	Aromatase immunoreactivity is increased in mammographically dense regions of the breast. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 243-252.	2.5	48
96	Extent of atypical hyperplasia stratifies breast cancer risk in 2 independent cohorts of women. <i>Cancer</i> , 2016, 122, 2971-2978.	4.1	48
97	The influence of mammogram acquisition on the mammographic density and breast cancer association in the mayo mammography health study cohort. <i>Breast Cancer Research</i> , 2012, 14, R147.	5.0	47
98	Differences in genomic abnormalities among African individuals with monoclonal gammopathies using calculated ancestry. <i>Blood Cancer Journal</i> , 2018, 8, 96.	6.2	47
99	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.	1.6	47
100	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	1.9	45
101	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
102	Mammographic density, parity and age at first birth, and risk of breast cancer: an analysis of four case-control studies. <i>Breast Cancer Research and Treatment</i> , 2012, 132, 1163-1171.	2.5	43
103	Prospective evaluation of a breast-cancer risk model integrating classical risk factors and polygenic risk in 15 cohorts from six countries. <i>International Journal of Epidemiology</i> , 2022, 50, 1897-1911.	1.9	43
104	Prenatal and Perinatal Correlates of Adult Mammographic Breast Density. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 1502-1508.	2.5	42
105	Dense and Nondense Mammographic Area and Risk of Breast Cancer by Age and Tumor Characteristics. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 798-809.	2.5	42
106	Genetic variation in stromal proteins decorin and lumican with breast cancer: investigations in two case-control studies. <i>Breast Cancer Research</i> , 2008, 10, R98.	5.0	41
107	Investigation of an interaction of alcohol intake and family history on breast cancer risk in the Minnesota Breast Cancer Family Study. <i>Cancer</i> , 2001, 92, 240-248.	4.1	40
108	Can genes for mammographic density inform cancer aetiology?. <i>Nature Reviews Cancer</i> , 2008, 8, 812-823.	28.4	40

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109	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	2.9	40
110	Case-control study of increased mammographic breast density response to hormone replacement therapy. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002, 11, 1382-8.	2.5	40
111	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	6.2	39
112	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. <i>Nature Communications</i> , 2015, 6, 7539.	12.8	38
113	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
114	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
115	Longitudinal Changes in Volumetric Breast Density with Tamoxifen and Aromatase Inhibitors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 930-937.	2.5	37
116	Breast Cancer Screening Strategies for Women With <i>ATM</i> , <i>CHEK2</i> , and <i>PALB2</i> Pathogenic Variants. <i>JAMA Oncology</i> , 2022, 8, 587.	7.1	36
117	Mammographic texture and risk of breast cancer by tumor type and estrogen receptor status. <i>Breast Cancer Research</i> , 2016, 18, 122.	5.0	35
118	Combining quantitative and qualitative breast density measures to assess breast cancer risk. <i>Breast Cancer Research</i> , 2017, 19, 97.	5.0	35
119	Association of Parity and Ovarian Cancer Risk by Family History of Breast or Ovarian Cancer in a Population-Based Study of Postmenopausal Women. <i>Epidemiology</i> , 2002, 13, 66-71.	2.7	34
120	Mammographic density does not differ between unaffected BRCA1/2 mutation carriers and women at low-to-average risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2010, 123, 245-255.	2.5	33
121	Comparison of percent density from raw and processed full-field digital mammography data. <i>Breast Cancer Research</i> , 2013, 15, R1.	5.0	32
122	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46 450 cases and 42 461 controls from the breast cancer association consortium. <i>Human Molecular Genetics</i> , 2014, 23, 1934-1946.	2.9	32
123	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	1.3	32
124	Identification of a novel percent mammographic density locus at 12q24. <i>Human Molecular Genetics</i> , 2012, 21, 3299-3305.	2.9	31
125	Mammographic texture resemblance generalizes as an independent risk factor for breast cancer. <i>Breast Cancer Research</i> , 2014, 16, R37.	5.0	31
126	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

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127	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
128	Mammographic density and risk of breast cancer by adiposity: An analysis of four case-control studies. <i>International Journal of Cancer</i> , 2012, 130, 1915-1924.	5.1	30
129	Mammographic Breast Density Response to Aromatase Inhibition. <i>Clinical Cancer Research</i> , 2013, 19, 2144-2153.	7.0	30
130	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	12.8	30
131	Novel pedigree analysis implicates DNA repair and chromatin remodeling in multiple myeloma risk. <i>PLoS Genetics</i> , 2018, 14, e1007111.	3.5	30
132	Natural history of age-related lobular involution and impact on breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 423-430.	2.5	29
133	Deep-LIBRA: An artificial-intelligence method for robust quantification of breast density with independent validation in breast cancer risk assessment. <i>Medical Image Analysis</i> , 2021, 73, 102138.	11.6	29
134	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
135	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	5.0	26
136	Whole Genome Sequence of Multiple Myeloma-Prone C57BL/KaLwRij Mouse Strain Suggests the Origin of Disease Involves Multiple Cell Types. <i>PLoS ONE</i> , 2015, 10, e0127828.	2.5	26
137	Breast Density Awareness, Knowledge, and Attitudes Among US Women: National Survey Results Across 5 Years. <i>Journal of the American College of Radiology</i> , 2020, 17, 391-404.	1.8	26
138	Association of diabetes with mammographic breast density and breast cancer in the Minnesota breast cancer family study. <i>Cancer Causes and Control</i> , 2007, 18, 505-515.	1.8	25
139	Mediterranean Diet and Breast Density in the Minnesota Breast Cancer Family Study. <i>Nutrition and Cancer</i> , 2008, 60, 703-709.	2.0	24
140	Background Parenchymal Uptake During Molecular Breast Imaging and Associated Clinical Factors. <i>American Journal of Roentgenology</i> , 2015, 204, W363-W370.	2.2	24
141	Association between mammographic breast density and histologic features of benign breast disease. <i>Breast Cancer Research</i> , 2017, 19, 134.	5.0	24
142	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.	5.0	24
143	Mammographic breast density and risk of breast cancer in women with atypical hyperplasia: an observational cohort study from the Mayo Clinic Benign Breast Disease (BBD) cohort. <i>BMC Cancer</i> , 2017, 17, 84.	2.6	23
144	Alcohol consumption and breast tumor gene expression. <i>Breast Cancer Research</i> , 2017, 19, 108.	5.0	23

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145	A polygenic risk score for breast cancer in women receiving tamoxifen or raloxifene on NSABP P-1 and P-2. <i>Breast Cancer Research and Treatment</i> , 2015, 149, 517-523.	2.5	22
146	Model for Predicting Breast Cancer Risk in Women With Atypical Hyperplasia. <i>Journal of Clinical Oncology</i> , 2018, 36, 1840-1846.	1.6	22
147	Fully Automated Volumetric Breast Density Estimation from Digital Breast Tomosynthesis. <i>Radiology</i> , 2021, 301, 561-568.	7.3	22
148	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.	1.6	22
149	Alcohol intake in adolescence and mammographic density. <i>International Journal of Cancer</i> , 2005, 117, 837-841.	5.1	21
150	Association of polygenic risk score with the risk of chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. <i>Blood</i> , 2018, 131, 2541-2551.	1.4	21
151	Evaluation of LIBRA Software for Fully Automated Mammographic Density Assessment in Breast Cancer Risk Prediction. <i>Radiology</i> , 2020, 296, 24-31.	7.3	21
152	Risk of Late-Onset Breast Cancer in Genetically Predisposed Women. <i>Journal of Clinical Oncology</i> , 2021, 39, 3430-3440.	1.6	21
153	Lobular involution: localized phenomenon or field effect?. <i>Breast Cancer Research and Treatment</i> , 2009, 117, 193-196.	2.5	20
154	Postmenopausal mammographic breast density and subsequent breast cancer risk according to selected tissue markers. <i>British Journal of Cancer</i> , 2015, 113, 1104-1113.	6.4	20
155	Personalizing Aspirin Use for Targeted Breast Cancer Chemoprevention in Postmenopausal Women. <i>Mayo Clinic Proceedings</i> , 2016, 91, 71-80.	3.0	20
156	Coinherited genetics of multiple myeloma and its precursor, monoclonal gammopathy of undetermined significance. <i>Blood Advances</i> , 2020, 4, 2789-2797.	5.2	20
157	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
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