

Celine M Vachon

List of Publications by Year in descending order

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Version: 2024-02-01

279
papers

20,392
citations

12303

69
h-index

12910

131
g-index

283
all docs

283
docs citations

283
times ranked

19510
citing authors

#	ARTICLE	IF	CITATIONS
1	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.	0.9	43
2	Polygenic risk score and risk of monoclonal B-cell lymphocytosis in caucasians and risk of chronic lymphocytic leukemia (CLL) in African Americans. <i>Leukemia</i> , 2022, 36, 119-125.	3.3	10
3	Family history of plasma cell disorders is associated with improved survival in MGUS, multiple myeloma, and systemic AL amyloidosis. <i>Leukemia</i> , 2022, 36, 1058-1065.	3.3	3
4	Evaluating educational interventions to increase breast density awareness among Latinas: A randomized trial in a Federally Qualified Health Center. <i>Cancer</i> , 2022, 128, 1038-1047.	2.0	5
5	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
6	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
7	Breast Density Knowledge and Awareness Among Latinas in a Low-Resource Setting. <i>Journal of the American College of Radiology</i> , 2022, 19, 155-161.	0.9	3
8	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.	3.4	36
9	Associations of history of vaccination and hospitalization due to infection with risk of monoclonal B-cell lymphocytosis. <i>Leukemia</i> , 2022, , .	3.3	1
10	Prevalence of heavy chain MGUS by race and family history risk groups using a high-sensitivity screening method. <i>Blood Advances</i> , 2022, 6, 3746-3750.	2.5	2
11	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
12	Body mass index associated with monoclonal gammopathy of undetermined significance (MGUS) progression in Olmsted County, Minnesota. <i>Blood Cancer Journal</i> , 2022, 12, 67.	2.8	13
13	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
14	Genome-wide meta-analysis of monoclonal gammopathy of undetermined significance (MGUS) identifies risk loci impacting IRF-6. <i>Blood Cancer Journal</i> , 2022, 12, 60.	2.8	2
15	Association of breast cancer risk, density, and stiffness: global tissue stiffness on breast MR elastography (MRE). <i>Breast Cancer Research and Treatment</i> , 2022, 194, 79-89.	1.1	9
16	Distinct Reproductive Risk Profiles for Intrinsic-Like Breast Cancer Subtypes: Pooled Analysis of Population-Based Studies. <i>Journal of the National Cancer Institute</i> , 2022, 114, 1706-1719.	3.0	14
17	Antimullerian Hormone as a Serum Biomarker for Risk of Chemotherapy-Induced Amenorrhea. <i>Journal of the National Cancer Institute</i> , 2021, 113, 1105-1108.	3.0	5
18	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

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19	Background Parenchymal Uptake on Molecular Breast Imaging and Breast Cancer Risk: A Cohort Study. <i>American Journal of Roentgenology</i> , 2021, 216, 1193-1204.	1.0	11
20	Impact of Personalized Genetic Breast Cancer Risk Estimation With Polygenic Risk Scores on Preventive Endocrine Therapy Intention and Uptake. <i>Cancer Prevention Research</i> , 2021, 14, 175-184.	0.7	11
21	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
22	Association of mammographic density measures and breast cancer intrinsic molecular subtypes. <i>Breast Cancer Research and Treatment</i> , 2021, 187, 215-224.	1.1	11
23	Genetic Variations and Health-Related Quality of Life (HRQOL): A Genome-Wide Study Approach. <i>Cancers</i> , 2021, 13, 716.	1.7	3
24	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
25	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
26	Association of Daily Alcohol Intake, Volumetric Breast Density, and Breast Cancer Risk. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkaa124.	1.4	2
27	Sequencing at lymphoid neoplasm susceptibility loci maps six myeloma risk genes. <i>Human Molecular Genetics</i> , 2021, 30, 1142-1153.	1.4	2
28	Genetic Predictors of Chemotherapy-Induced Peripheral Neuropathy from Paclitaxel, Carboplatin and Oxaliplatin: NCCTG/Alliance N08C1, N08CA and N08CB Study. <i>Cancers</i> , 2021, 13, 1084.	1.7	11
29	Expression quantitative trait loci of genes predicting outcome are associated with survival of multiple myeloma patients. <i>International Journal of Cancer</i> , 2021, 149, 327-336.	2.3	3
30	Natural history of monoclonal B-cell lymphocytosis among relatives in CLL families. <i>Blood</i> , 2021, 137, 2046-2056.	0.6	16
31	Automated Quantitative Measures of Terminal Duct Lobular Unit Involution and Breast Cancer Risk Letter. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 797-797.	1.1	1
32	Automated percent mammographic density, mammographic texture variation, and risk of breast cancer: a nested case-control study. <i>Npj Breast Cancer</i> , 2021, 7, 68.	2.3	15
33	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
34	Risk of Late-Onset Breast Cancer in Genetically Predisposed Women. <i>Journal of Clinical Oncology</i> , 2021, 39, 3430-3440.	0.8	21
35	Mammographic Variation Measures, Breast Density, and Breast Cancer Risk. <i>American Journal of Roentgenology</i> , 2021, 217, 326-335.	1.0	9
36	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

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37	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
38	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
39	Genetic determinants of multiple myeloma risk within the Wnt/beta-catenin signaling pathway. <i>Cancer Epidemiology</i> , 2021, 73, 101972.	0.8	0
40	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
41	Deep Learning Predicts Interval and Screening-detected Cancer from Screening Mammograms: A Case-Case-Control Study in 6369 Women. <i>Radiology</i> , 2021, 301, 550-558.	3.6	15
42	Fully Automated Volumetric Breast Density Estimation from Digital Breast Tomosynthesis. <i>Radiology</i> , 2021, 301, 561-568.	3.6	22
43	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.	7.0	29
44	Factors Associated With Severe COVID-19 Infection Among Persons of Different Ages Living in a Defined Midwestern US Population. <i>Mayo Clinic Proceedings</i> , 2021, 96, 2528-2539.	1.4	16
45	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
46	Simplified Breast Risk Tool Integrating Questionnaire Risk Factors, Mammographic Density, and Polygenic Risk Score: Development and Validation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 600-607.	1.1	14
47	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
48	Association of EDARV370A with breast density and metabolic syndrome in Latinos. <i>PLoS ONE</i> , 2021, 16, e0258212.	1.1	5
49	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
50	Incorporating Robustness to Imaging Physics into Radiomic Feature Selection for Breast Cancer Risk Estimation. <i>Cancers</i> , 2021, 13, 5497.	1.7	4
51	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
52	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.	0.9	26
53	Breast Cancer Risk and Use of Nonsteroidal Anti-inflammatory Agents After a Benign Breast Biopsy. <i>Cancer Prevention Research</i> , 2020, 13, 967-976.	0.7	9
54	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	2.6	39

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55	Free Light Chain Assay Drift: Potential for Misdiagnosis?. <i>Journal of applied laboratory medicine</i> , The, 2020, 5, 1411-1413.	0.6	15
56	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
57	Evaluation of LIBRA Software for Fully Automated Mammographic Density Assessment in Breast Cancer Risk Prediction. <i>Radiology</i> , 2020, 296, 24-31.	3.6	21
58	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
59	The CCND1 c.870G risk allele is enriched in individuals of African ancestry with plasma cell dyscrasias. <i>Blood Cancer Journal</i> , 2020, 10, 39.	2.8	4
60	Coinherited genetics of multiple myeloma and its precursor, monoclonal gammopathy of undetermined significance. <i>Blood Advances</i> , 2020, 4, 2789-2797.	2.5	20
61	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
62	The Association of Modifiable Breast Cancer Risk Factors and Somatic Genomic Alterations in Breast Tumors: The Cancer Genome Atlas Network. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 599-605.	1.1	7
63	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
64	Tumor mutational load predicts time to first treatment in chronic lymphocytic leukemia (CLL) and monoclonal B-cell lymphocytosis beyond the CLL international prognostic index. <i>American Journal of Hematology</i> , 2020, 95, 906-917.	2.0	17
65	Assessing Vitamin D and Mammographic Breast Density in Alaskan Women. <i>Clinics and Practice</i> , 2020, 10, 1253.	0.6	3
66	Polygenic Risk Score and Risk of Chronic Lymphocytic Leukemia, Monoclonal B-Cell Lymphocytosis (MBL), and MBL Subtypes. <i>Blood</i> , 2020, 136, 35-36.	0.6	0
67	Comparison of MGUS Prevalence By Race and Family History Risk Groups Using a High Sensitivity Screening Method (MASS-FIX). <i>Blood</i> , 2020, 136, 40-41.	0.6	1
68	Body Mass Index and Clinical Factors Associated with Monoclonal Gammopathy of Undetermined Significance (MGUS) Progression in Olmsted County, Minnesota. <i>Blood</i> , 2020, 136, 15-16.	0.6	0
69	Prevalence of Familial Plasma Cell Disorders in Patients with Multiple Myeloma. <i>Blood</i> , 2020, 136, 1-2.	0.6	0
70	Association of elevated serumfree light chains with chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. <i>Blood Cancer Journal</i> , 2019, 9, 59.	2.8	9
71	Genetic predictors of chemotherapy-related amenorrhea in women with breast cancer. <i>Fertility and Sterility</i> , 2019, 112, 731-739.e1.	0.5	10
72	Associations of mammographic breast density with breast stem cell marker-defined breast cancer subtypes. <i>Cancer Causes and Control</i> , 2019, 30, 1103-1111.	0.8	7

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73	Automated volumetric breast density measures: differential change between breasts in women with and without breast cancer. <i>Breast Cancer Research</i> , 2019, 21, 118.	2.2	13
74	Accuracy of self-reported cancer treatment data in young breast cancer survivors. <i>Journal of Patient-Reported Outcomes</i> , 2019, 3, 24.	0.9	5
75	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
76	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
77	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.	2.2	31
78	Discussions of Dense Breasts, Breast Cancer Risk, and Screening Choices in 2019. <i>JAMA - Journal of the American Medical Association</i> , 2019, 322, 69.	3.8	15
79	Longitudinal Changes in Volumetric Breast Density in Healthy Women across the Menopausal Transition. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1324-1330.	1.1	17
80	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
81	Impact of short-term low-dose tamoxifen on molecular breast imaging background parenchymal uptake: a pilot study. <i>Breast Cancer Research</i> , 2019, 21, 38.	2.2	11
82	Body mass index, mammographic density, and breast cancer risk by estrogen receptor subtype. <i>Breast Cancer Research</i> , 2019, 21, 48.	2.2	52
83	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
84	Incidence of AL Amyloidosis in Olmsted County, Minnesota, 1990 through 2015. <i>Mayo Clinic Proceedings</i> , 2019, 94, 465-471.	1.4	87
85	Breast Cancer Classification using Deep Transfer Learning on Structured Healthcare Data. , 2019, , .		9
86	Behavioral and psychological impact of returning breast density results to Latinas: study protocol for a randomized clinical trial. <i>Trials</i> , 2019, 20, 744.	0.7	5
87	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.	2.8	57
88	Risk of MGUS in relatives of multiple myeloma cases by clinical and tumor characteristics. <i>Leukemia</i> , 2019, 33, 499-507.	3.3	9
89	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
90	Molecular mechanisms linking high body mass index to breast cancer etiology in post-menopausal breast tumor and tumor-adjacent tissues. <i>Breast Cancer Research and Treatment</i> , 2019, 173, 667-677.	1.1	19

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91	Radiomic Phenotypes of Mammographic Parenchymal Complexity: Toward Augmenting Breast Density in Breast Cancer Risk Assessment. <i>Radiology</i> , 2019, 290, 41-49.	3.6	63
92	Association between a Polygenic Risk Score for Multiple Myeloma Risk and Overall Survival. <i>Blood</i> , 2019, 134, 4366-4366.	0.6	0
93	Germline Variation Predicts Treatment Response in Multiple Myeloma. <i>Blood</i> , 2019, 134, 4397-4397.	0.6	0
94	The CCND1 870G Risk Allele Is Enriched in African Individuals with Plasma Cell Dyscrasias. <i>Blood</i> , 2019, 134, 4362-4362.	0.6	0
95	Tumor Mutational Load and Germline Polygenic Risk Score Predicts Time-to-First Treatment in Chronic Lymphocytic Leukemia (CLL) and High-Count Monoclonal B Cell Lymphocytosis (MBL). <i>Blood</i> , 2019, 134, 852-852.	0.6	0
96	Stroma modifies relationships between risk factor exposure and age-related epithelial involution in benign breast. <i>Modern Pathology</i> , 2018, 31, 1085-1096.	2.9	9
97	Does mammographic density mediate risk factor associations with breast cancer? An analysis by tumor characteristics. <i>Breast Cancer Research and Treatment</i> , 2018, 170, 129-141.	1.1	11
98	Association of polygenic risk score with the risk of chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. <i>Blood</i> , 2018, 131, 2541-2551.	0.6	21
99	Prenatal diethylstilbestrol exposure and mammographic density. <i>International Journal of Cancer</i> , 2018, 143, 1374-1378.	2.3	3
100	Common Genetic Variation and Breast Cancer Risk—Past, Present, and Future. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 380-394.	1.1	108
101	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.	0.9	88
102	Germline Lysine-Specific Demethylase 1 (<i>LSD1/KDM1A</i>) Mutations Confer Susceptibility to Multiple Myeloma. <i>Cancer Research</i> , 2018, 78, 2747-2759.	0.4	56
103	Model for Predicting Breast Cancer Risk in Women With Atypical Hyperplasia. <i>Journal of Clinical Oncology</i> , 2018, 36, 1840-1846.	0.8	22
104	Differences in genomic abnormalities among African individuals with monoclonal gammopathies using calculated ancestry. <i>Blood Cancer Journal</i> , 2018, 8, 96.	2.8	47
105	Evaluation of 2 breast cancer risk models in a benign breast disease cohort. <i>Cancer</i> , 2018, 124, 3319-3328.	2.0	7
106	Quantitative background parenchymal uptake on molecular breast imaging and breast cancer risk: a case-control study. <i>Breast Cancer Research</i> , 2018, 20, 46.	2.2	12
107	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.	2.0	56
108	The National Cancer Institute Cohort Consortium: An International Pooling Collaboration of 58 Cohorts from 20 Countries. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 1307-1319.	1.1	18

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109	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
110	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
111	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
112	Polygenic risk score for breast cancer in high-risk women.. <i>Journal of Clinical Oncology</i> , 2018, 36, 1508-1508.	0.8	11
113	Novel pedigree analysis implicates DNA repair and chromatin remodeling in multiple myeloma risk. <i>PLoS Genetics</i> , 2018, 14, e1007111.	1.5	30
114	Accuracy of self-reported chemotherapy regimens in young breast cancer survivors.. <i>Journal of Clinical Oncology</i> , 2018, 36, e22143-e22143.	0.8	0
115	Clonal Hematopoiesis of Indeterminate Potential (CHIP) and Chronic Lymphocytic Leukemia (CLL) Driver Genes: Risk of CLL and Monoclonal B-Cell Lymphocytosis (MBL). <i>Blood</i> , 2018, 132, 3116-3116.	0.6	0
116	Genomic Abnormalities Among African Individuals with Monoclonal Gammopathies Using Calculated Ancestry. <i>Blood</i> , 2018, 132, 4458-4458.	0.6	0
117	Large-Scale Linkage Analysis of Multiple Myeloma (MM) and Monoclonal Gammopathy of Undetermined Significance (MGUS) Families. <i>Blood</i> , 2018, 132, 4501-4501.	0.6	0
118	Longitudinal Changes in Volumetric Breast Density with Tamoxifen and Aromatase Inhibitors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 930-937.	1.1	37
119	Interaction of mammographic breast density with menopausal status and postmenopausal hormone use in relation to the risk of aggressive breast cancer subtypes. <i>Breast Cancer Research and Treatment</i> , 2017, 165, 421-431.	1.1	11
120	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.	1.1	23
121	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
122	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
123	NanoString-based breast cancer risk prediction for women with sclerosing adenosis. <i>Breast Cancer Research and Treatment</i> , 2017, 166, 641-650.	1.1	10
124	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	45
125	Combining quantitative and qualitative breast density measures to assess breast cancer risk. <i>Breast Cancer Research</i> , 2017, 19, 97.	2.2	35
126	Tissue-based associations of mammographic breast density with breast stem cell markers. <i>Breast Cancer Research</i> , 2017, 19, 100.	2.2	7

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127	Alcohol consumption and breast tumor gene expression. <i>Breast Cancer Research</i> , 2017, 19, 108.	2.2	23
128	Association between mammographic breast density and histologic features of benign breast disease. <i>Breast Cancer Research</i> , 2017, 19, 134.	2.2	24
129	Mammographic density and ageing: A collaborative pooled analysis of cross-sectional data from 22 countries worldwide. <i>PLoS Medicine</i> , 2017, 14, e1002335.	3.9	108
130	Using Breast Cancer Risk Associated Polymorphisms to Identify Women for Breast Cancer Chemoprevention. <i>PLoS ONE</i> , 2017, 12, e0168601.	1.1	16
131	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
132	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	1.1	12
133	<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
134	Mammographic texture and risk of breast cancer by tumor type and estrogen receptor status. <i>Breast Cancer Research</i> , 2016, 18, 122.	2.2	35
135	Unsupervised Deep Learning Applied to Breast Density Segmentation and Mammographic Risk Scoring. <i>IEEE Transactions on Medical Imaging</i> , 2016, 35, 1322-1331.	5.4	360
136	Genes associated with histopathologic features of triple negative breast tumors predict molecular subtypes. <i>Breast Cancer Research and Treatment</i> , 2016, 157, 117-131.	1.1	18
137	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1609-1618.	1.1	18
138	Breast cancer risk by the extent and type of atypical hyperplasia. <i>Cancer</i> , 2016, 122, 3087-3088.	2.0	10
139	Extent of atypical hyperplasia stratifies breast cancer risk in 2 independent cohorts of women. <i>Cancer</i> , 2016, 122, 2971-2978.	2.0	48
140	Breast cancer risk prediction using a clinical risk model and polygenic risk score. <i>Breast Cancer Research and Treatment</i> , 2016, 159, 513-525.	1.1	129
141	rs2735383, located at a microRNA binding site in the 3'UTR of NBS1, is not associated with breast cancer risk. <i>Scientific Reports</i> , 2016, 6, 36874.	1.6	2
142	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
143	Mammographic density is the main correlate of tumors detected on ultrasound but not on mammography. <i>International Journal of Cancer</i> , 2016, 139, 1967-1974.	2.3	19
144	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93

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145	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
146	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
147	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	5.8	94
148	Mammographic density assessed on paired raw and processed digital images and on paired screen-film and digital images across three mammography systems. <i>Breast Cancer Research</i> , 2016, 18, 130.	2.2	17
149	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for <i>CHK2</i> *1100delC Carriers. <i>Journal of Clinical Oncology</i> , 2016, 34, 2750-2760.	0.8	152
150	Breast Density: More Than Meets the Eye. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw128.	3.0	6
151	Background parenchymal uptake on molecular breast imaging as a breast cancer risk factor: a case-control study. <i>Breast Cancer Research</i> , 2016, 18, 42.	2.2	19
152	International Consortium on Mammographic Density: Methodology and population diversity captured across 22 countries. <i>Cancer Epidemiology</i> , 2016, 40, 141-151.	0.8	19
153	Personalizing Aspirin Use for Targeted Breast Cancer Chemoprevention in Postmenopausal Women. <i>Mayo Clinic Proceedings</i> , 2016, 91, 71-80.	1.4	20
154	Natural history of age-related lobular involution and impact on breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 423-430.	1.1	29
155	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
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