

# 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	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
3	Benign Breast Disease and the Risk of Breast Cancer. <i>New England Journal of Medicine</i> , 2005, 353, 229-237.	13.9	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.	2.6	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.	3.0	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.	0.8	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.	9.4	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.	9.4	493
9	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	428
10	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
11	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	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.	5.4	360
13	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	1.5	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.	0.8	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.	9.4	309
16	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	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.	9.4	279
18	Mammographic density, breast cancer risk and risk prediction. <i>Breast Cancer Research</i> , 2007, 9, 217.	2.2	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.	9.4	265
20	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. <i>Journal of the National Cancer Institute</i> , 2014, 106, .	3.0	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.	3.0	226
22	Age-Related Lobular Involution and Risk of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2006, 98, 1600-1607.	3.0	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.	3.0	192
24	Breast Cancer Risk by Breast Density, Menopause, and Postmenopausal Hormone Therapy Use. <i>Journal of Clinical Oncology</i> , 2010, 28, 3830-3837.	0.8	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.	9.4	184
26	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
27	Mammographic Breast Density as a General Marker of Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 43-49.	1.1	181
28	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876.	9.4	179
29	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	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, .	3.0	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.	1.5	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.	0.8	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.	1.4	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.	7.7	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.	0.8	152
36	Mammographic density and risk of breast cancer by age and tumor characteristics. <i>Breast Cancer Research</i> , 2013, 15, R104.	2.2	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.	1.3	145
38	Comparison of Clinical and Automated Breast Density Measurements: Implications for Risk Prediction and Supplemental Screening. <i>Radiology</i> , 2016, 279, 710-719.	3.6	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.	1.1	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.	0.6	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.	9.4	125
42	Texture Features from Mammographic Images and Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 837-845.	1.1	121
43	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
44	Longitudinal Trends in Mammographic Percent Density and Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 921-928.	1.1	118
45	Genome-wide association study identifies a novel susceptibility locus at 6p21.3 among familial CLL. <i>Blood</i> , 2011, 117, 1911-1916.	0.6	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.	1.8	118
47	Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. <i>Nature Genetics</i> , 2011, 43, 185-187.	9.4	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.	5.8	109
49	Common Genetic Variation and Breast Cancer Risk—Past, Present, and Future. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 380-394.	1.1	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.	3.9	108
51	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	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.	1.1	101
53	19p13.1 Is a Triple-Negative-Specific Breast Cancer Susceptibility Locus. <i>Cancer Research</i> , 2012, 72, 1795-1803.	0.4	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.	3.0	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.	3.0	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.	2.6	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.	0.8	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.	0.8	94
59	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	5.8	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.	1.5	94
61	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
62	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
63	Age-specific Trends in Mammographic Density: The Minnesota Breast Cancer Family Study. <i>American Journal of Epidemiology</i> , 2008, 167, 1027-1036.	1.6	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.	0.9	88
65	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
66	Incidence of AL Amyloidosis in Olmsted County, Minnesota, 1990 through 2015. <i>Mayo Clinic Proceedings</i> , 2019, 94, 465-471.	1.4	87
67	Association Between Mammographic Density and Age-Related Lobular Involution of the Breast. <i>Journal of Clinical Oncology</i> , 2010, 28, 2207-2212.	0.8	84
68	Breast Density and Breast Cancer Risk: A Practical Review. <i>Mayo Clinic Proceedings</i> , 2014, 89, 548-557.	1.4	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.	1.1	83
70	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
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.	2.6	76
72	Tissue composition of mammographically dense and non-dense breast tissue. <i>Breast Cancer Research and Treatment</i> , 2012, 131, 267-275.	1.1	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.	0.4	69
74	Familial monoclonal gammopathy of undetermined significance and multiple myeloma: epidemiology, risk factors, and biological characteristics. <i>Blood</i> , 2012, 119, 5359-5366.	0.6	68
75	Association of Genetic Variation in Genes Implicated in the $\beta$ -Catenin Destruction Complex with Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2101-2108.	1.1	67
76	An Automated Approach for Estimation of Breast Density. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3090-3097.	1.1	67
77	A Novel Automated Mammographic Density Measure and Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2012, 104, 1028-1037.	3.0	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.	3.0	66
79	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
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.	1.2	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.	1.1	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.	2.8	57
83	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	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.	0.4	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.	2.0	56
86	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , 2015, 75, 2457-2467.	0.4	55
87	Mammographic Breast Density and Breast Cancer: Evidence of a Shared Genetic Basis. <i>Cancer Research</i> , 2012, 72, 1478-1484.	0.4	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.	1.4	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.	1.1	52
90	Body mass index, mammographic density, and breast cancer risk by estrogen receptor subtype. <i>Breast Cancer Research</i> , 2019, 21, 48.	2.2	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.	2.9	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.	0.8	51
93	Annexin A1 expression in a pooled breast cancer series: association with tumor subtypes and prognosis. <i>BMC Medicine</i> , 2015, 13, 156.	2.3	51
94	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	1.1	49
95	Aromatase immunoreactivity is increased in mammographically dense regions of the breast. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 243-252.	1.1	48
96	Extent of atypical hyperplasia stratifies breast cancer risk in 2 independent cohorts of women. <i>Cancer</i> , 2016, 122, 2971-2978.	2.0	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.	2.2	47
98	Differences in genomic abnormalities among African individuals with monoclonal gammopathies using calculated ancestry. <i>Blood Cancer Journal</i> , 2018, 8, 96.	2.8	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.	0.8	47
100	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	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.	3.0	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.	1.1	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.	0.9	43
104	Prenatal and Perinatal Correlates of Adult Mammographic Breast Density. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 1502-1508.	1.1	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.	1.1	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.	2.2	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.	2.0	40
108	Can genes for mammographic density inform cancer aetiology?. <i>Nature Reviews Cancer</i> , 2008, 8, 812-823.	12.8	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.	1.4	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.	1.1	40
111	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	2.6	39
112	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. <i>Nature Communications</i> , 2015, 6, 7539.	5.8	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.	1.4	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.	2.6	37
115	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
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.	3.4	36
117	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
118	Combining quantitative and qualitative breast density measures to assess breast cancer risk. <i>Breast Cancer Research</i> , 2017, 19, 97.	2.2	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.	1.2	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.	1.1	33
121	Comparison of percent density from raw and processed full-field digital mammography data. <i>Breast Cancer Research</i> , 2013, 15, R1.	2.2	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.	1.4	32
123	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
124	Identification of a novel percent mammographic density locus at 12q24. <i>Human Molecular Genetics</i> , 2012, 21, 3299-3305.	1.4	31
125	Mammographic texture resemblance generalizes as an independent risk factor for breast cancer. <i>Breast Cancer Research</i> , 2014, 16, R37.	2.2	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.	0.8	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.	2.2	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.	2.3	30
129	Mammographic Breast Density Response to Aromatase Inhibition. <i>Clinical Cancer Research</i> , 2013, 19, 2144-2153.	3.2	30
130	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
131	Novel pedigree analysis implicates DNA repair and chromatin remodeling in multiple myeloma risk. <i>PLoS Genetics</i> , 2018, 14, e1007111.	1.5	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.	1.1	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.	7.0	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.	1.6	26
135	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	2.2	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.	1.1	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.	0.9	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.	0.8	25
139	Mediterranean Diet and Breast Density in the Minnesota Breast Cancer Family Study. <i>Nutrition and Cancer</i> , 2008, 60, 703-709.	0.9	24
140	Background Parenchymal Uptake During Molecular Breast Imaging and Associated Clinical Factors. <i>American Journal of Roentgenology</i> , 2015, 204, W363-W370.	1.0	24
141	Association between mammographic breast density and histologic features of benign breast disease. <i>Breast Cancer Research</i> , 2017, 19, 134.	2.2	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.	2.2	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.	1.1	23
144	Alcohol consumption and breast tumor gene expression. <i>Breast Cancer Research</i> , 2017, 19, 108.	2.2	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.	1.1	22
146	Model for Predicting Breast Cancer Risk in Women With Atypical Hyperplasia. <i>Journal of Clinical Oncology</i> , 2018, 36, 1840-1846.	0.8	22
147	Fully Automated Volumetric Breast Density Estimation from Digital Breast Tomosynthesis. <i>Radiology</i> , 2021, 301, 561-568.	3.6	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.	0.8	22
149	Alcohol intake in adolescence and mammographic density. <i>International Journal of Cancer</i> , 2005, 117, 837-841.	2.3	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.	0.6	21
151	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
152	Risk of Late-Onset Breast Cancer in Genetically Predisposed Women. <i>Journal of Clinical Oncology</i> , 2021, 39, 3430-3440.	0.8	21
153	Lobular involution: localized phenomenon or field effect?. <i>Breast Cancer Research and Treatment</i> , 2009, 117, 193-196.	1.1	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.	2.9	20
155	Personalizing Aspirin Use for Targeted Breast Cancer Chemoprevention in Postmenopausal Women. <i>Mayo Clinic Proceedings</i> , 2016, 91, 71-80.	1.4	20
156	Coinherited genetics of multiple myeloma and its precursor, monoclonal gammopathy of undetermined significance. <i>Blood Advances</i> , 2020, 4, 2789-2797.	2.5	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.	2.2	19
158	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
159	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
160	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
161	International Consortium on Mammographic Density: Methodology and population diversity captured across 22 countries. <i>Cancer Epidemiology</i> , 2016, 40, 141-151.	0.8	19
162	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

#	ARTICLE	IF	CITATIONS
163	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
164	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
165	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
166	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
167	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
168	Dietary patterns and breast density in the Minnesota Breast Cancer Family Study. <i>Cancer Causes and Control</i> , 2008, 19, 481-489.	0.8	17
169	Mammographic density and breast cancer risk by family history in women of white and Asian ancestry. <i>Cancer Causes and Control</i> , 2015, 26, 621-626.	0.8	17
170	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
171	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
172	Tumor mutational load predicts time to first treatment in chronic lymphocytic leukemia ( CLL ) and monoclonal Bâ€cell lymphocytosis beyond the CLL international prognostic index. <i>American Journal of Hematology</i> , 2020, 95, 906-917.	2.0	17
173	Genetic variation in the estrogen metabolic pathway and mammographic density as an intermediate phenotype of breast cancer. <i>Breast Cancer Research</i> , 2010, 12, R19.	2.2	16
174	Natural history of monoclonal B-cell lymphocytosis among relatives in CLL families. <i>Blood</i> , 2021, 137, 2046-2056.	0.6	16
175	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
176	Using Breast Cancer Risk Associated Polymorphisms to Identify Women for Breast Cancer Chemoprevention. <i>PLoS ONE</i> , 2017, 12, e0168601.	1.1	16
177	Automated Percentage of Breast Density Measurements for Full-field Digital Mammography Applications. <i>Academic Radiology</i> , 2014, 21, 958-970.	1.3	15
178	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
179	Free Light Chain Assay Drift: Potential for Misdiagnosis?. <i>journal of applied laboratory medicine</i> , The, 2020, 5, 1411-1413.	0.6	15
180	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

#	ARTICLE	IF	CITATIONS
181	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
182	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
183	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
184	Genetic anticipation and breast cancer: a prospective follow-up study. <i>Breast Cancer Research and Treatment</i> , 1999, 55, 21-28.	1.1	14
185	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. <i>Breast Cancer Research</i> , 2014, 16, R51.	2.2	14
186	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. <i>Carcinogenesis</i> , 2015, 36, 256-271.	1.3	14
187	Effect of Menstrual Cycle Phase on Background Parenchymal Uptake at Molecular Breast Imaging. <i>Academic Radiology</i> , 2015, 22, 1147-1156.	1.3	14
188	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
189	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
190	Familial correlation of dietary intakes among postmenopausal women. , 1998, 15, 553-563.		13
191	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
192	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
193	Increased prevalence of light chain monoclonal gammopathy of undetermined significance (<sc>LC</sc>â€<sc>MGUS</sc>) in first-degree relatives of individuals with multiple myeloma. <i>British Journal of Haematology</i> , 2012, 157, 472-475.	1.2	12
194	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	1.1	12
195	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
196	Evaluation of potential sources of bias in a genetic epidemiologic study of breast cancer. , 1997, 14, 85-95.		11
197	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
198	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

#	ARTICLE	IF	CITATIONS
199	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
200	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
201	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
202	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
203	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
204	Polygenic risk score for breast cancer in high-risk women.. <i>Journal of Clinical Oncology</i> , 2018, 36, 1508-1508.	0.8	11
205	Breast cancer risk by the extent and type of atypical hyperplasia. <i>Cancer</i> , 2016, 122, 3087-3088.	2.0	10
206	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
207	Genetic predictors of chemotherapy-related amenorrhea in women with breast cancer. <i>Fertility and Sterility</i> , 2019, 112, 731-739.e1.	0.5	10
208	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
209	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
210	Association of elevated serum free light chains with chronic lymphocytic leukemia and monoclonal B-cell lymphocytosis. <i>Blood Cancer Journal</i> , 2019, 9, 59.	2.8	9
211	Breast Cancer Classification using Deep Transfer Learning on Structured Healthcare Data. , 2019, , .		9
212	Risk of MGUS in relatives of multiple myeloma cases by clinical and tumor characteristics. <i>Leukemia</i> , 2019, 33, 499-507.	3.3	9
213	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
214	Mammographic Variation Measures, Breast Density, and Breast Cancer Risk. <i>American Journal of Roentgenology</i> , 2021, 217, 326-335.	1.0	9
215	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
216	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

#	ARTICLE	IF	CITATIONS
217	Methods for Assessing and Representing Mammographic Density: An Analysis of 4 Case-Control Studies. <i>American Journal of Epidemiology</i> , 2014, 179, 236-244.	1.6	8
218	Tissue-based associations of mammographic breast density with breast stem cell markers. <i>Breast Cancer Research</i> , 2017, 19, 100.	2.2	7
219	Evaluation of 2 breast cancer risk models in a benign breast disease cohort. <i>Cancer</i> , 2018, 124, 3319-3328.	2.0	7
220	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
221	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
222	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
223	Mammographic Density: Potential as a Risk Factor and Surrogate Marker in the Clinical Setting. <i>Current Breast Cancer Reports</i> , 2013, 5, 183-193.	0.5	6
224	Breast Density: More Than Meets the Eye. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw128.	3.0	6
225	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
226	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
227	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
228	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
229	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
230	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
231	CYP3A7*1C allele: linking premenopausal oestrogen and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	2.9	5
232	Association of EDARV370A with breast density and metabolic syndrome in Latinos. <i>PLoS ONE</i> , 2021, 16, e0258212.	1.1	5
233	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
234	Segregation analysis of breast cancer: A comparison of type-dependent age-at-onset versus type-dependent susceptibility models. , 1996, 13, 317-328.		4

#	ARTICLE	IF	CITATIONS
235	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
236	Skin Cancers Among Chronic Lymphocytic Leukemia (CLL) Patients - the Effect of UV Radiation and CLL Clinical Characteristics. <i>Blood</i> , 2016, 128, 4772-4772.	0.6	4
237	Incorporating Robustness to Imaging Physics into Radiomic Feature Selection for Breast Cancer Risk Estimation. <i>Cancers</i> , 2021, 13, 5497.	1.7	4
238	An investigation of the effects of mammographic acquisition parameters on a semiautomated quantitative measure of breast cancer risk. <i>Journal of Digital Imaging</i> , 2000, 13, 186-188.	1.6	3
239	Clinical Characteristics of Familial vs. Sporadic non-Hodgkin lymphoma in Patients Diagnosed at The Mayo Clinic (1986-2000). <i>Leukemia and Lymphoma</i> , 2004, 45, 929-935.	0.6	3
240	No evidence for association of inherited variation in genes involved in mitosis and percent mammographic density. <i>Breast Cancer Research</i> , 2012, 14, R7.	2.2	3
241	Prenatal diethylstilbestrol exposure and mammographic density. <i>International Journal of Cancer</i> , 2018, 143, 1374-1378.	2.3	3
242	Genetic Variations and Health-Related Quality of Life (HRQOL): A Genome-Wide Study Approach. <i>Cancers</i> , 2021, 13, 716.	1.7	3
243	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
244	Assessing Vitamin D and Mammographic Breast Density in Alaskan Women. <i>Clinics and Practice</i> , 2020, 10, 1253.	0.6	3
245	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
246	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
247	Differences in the distribution of cytogenetic subtypes between multiple myeloma patients with and without a family history of monoclonal gammopathy and multiple myeloma. <i>European Journal of Haematology</i> , 2013, 91, 193-195.	1.1	2
248	The association of copy number variation and percent mammographic density. <i>BMC Research Notes</i> , 2015, 8, 297.	0.6	2
249	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
250	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
251	Association of Daily Alcohol Intake, Volumetric Breast Density, and Breast Cancer Risk. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkaa124.	1.4	2
252	Sequencing at lymphoid neoplasm susceptibility loci maps six myeloma risk genes. <i>Human Molecular Genetics</i> , 2021, 30, 1142-1153.	1.4	2

#	ARTICLE	IF	CITATIONS
253	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
254	A Meta-Analysis Of Genome-Wide Association Studies Of Multiple Myeloma In Cases and Controls Of European Origin Identifies a Risk Locus In 12q23.1. <i>Blood</i> , 2013, 122, 3111-3111.	0.6	2
255	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
256	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
257	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
258	Does a Multiple Myeloma Polygenic Risk Score Predict Overall Survival of Myeloma Patients?. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 0, , .	1.1	2
259	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
260	Deletion Of Samsn1 Underlies Genetic Susceptibility To Monoclonal Gammopathy Of Undetermined Significance (MGUS) In Mice. <i>Blood</i> , 2013, 122, 397-397.	0.6	1
261	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
262	Associations of history of vaccination and hospitalization due to infection with risk of monoclonal B-cell lymphocytosis. <i>Leukemia</i> , 2022, , .	3.3	1
263	Response: Re: Age-Related Lobular Involution and Risk of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2007, 99, 572-572.	3.0	0
264	Genetic determinants of multiple myeloma risk within the Wnt/beta-catenin signaling pathway. <i>Cancer Epidemiology</i> , 2021, 73, 101972.	0.8	0
265	Increased Risk of Monoclonal Gammopathy in First-Degree Relatives of Patients with Multiple Myeloma or Monoclonal Gammopathy of Undetermined Significance.. <i>Blood</i> , 2008, 112, 1672-1672.	0.6	0
266	Prevalence of MBL Increases Over Time In Relatives of CLL Families,. <i>Blood</i> , 2011, 118, 3881-3881.	0.6	0
267	Differences in the Distribution of Cytogenetic Subtypes Between Multiple Myeloma Patients with and without a History of Familial MGUS and Multiple Myeloma. <i>Blood</i> , 2012, 120, 4000-4000.	0.6	0
268	Risk of Monoclonal Gammopathy of Undetermined Significance in First-Degree Relatives of Multiple Myeloma Cases By Cytogenetic Subtype. <i>Blood</i> , 2016, 128, 4425-4425.	0.6	0
269	Accuracy of self-reported chemotherapy regimens in young breast cancer survivors.. <i>Journal of Clinical Oncology</i> , 2018, 36, e22143-e22143.	0.8	0
270	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



#	ARTICLE	IF	CITATIONS
271	Genomic Abnormalities Among African Individuals with Monoclonal Gammopathies Using Calculated Ancestry. <i>Blood</i> , 2018, 132, 4458-4458.	0.6	0
272	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
273	Association between a Polygenic Risk Score for Multiple Myeloma Risk and Overall Survival. <i>Blood</i> , 2019, 134, 4366-4366.	0.6	0
274	Germline Variation Predicts Treatment Response in Multiple Myeloma. <i>Blood</i> , 2019, 134, 4397-4397.	0.6	0
275	The CCND1 870G Risk Allele Is Enriched in African Individuals with Plasma Cell Dyscrasias. <i>Blood</i> , 2019, 134, 4362-4362.	0.6	0
276	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
277	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
278	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
279	Prevalence of Familial Plasma Cell Disorders in Patients with Multiple Myeloma. <i>Blood</i> , 2020, 136, 1-2.	0.6	0