

# Jingmei Li

## List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

167  
papers

8,416  
citations

42  
h-index

88  
g-index

181  
ext. papers

10,557  
ext. citations

9.3  
avg, IF

4.51  
L-index

#	Paper	IF	Citations
167	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , <b>2013</b> , 45, 353-61, 361e1-2	36.3	813
166	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94	50.4	643
165	GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. <i>Science</i> , <b>2013</b> , 340, 1467-71	33.3	563
164	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , <b>2013</b> , 45, 371-84, 384e1-2	36.3	422
163	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 373-80	36.3	406
162	Prediction of breast cancer risk based on profiling with common genetic variants. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	324
161	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , <b>2017</b> , 49, 834-841	36.3	257
160	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , <b>2015</b> , 47, 1294-1303	36.3	226
159	Mammographic density phenotypes and risk of breast cancer: a meta-analysis. <i>Journal of the National Cancer Institute</i> , <b>2014</b> , 106,	9.7	190
158	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 1767-1778	36.3	186
157	Functional variants at the 11q13 risk locus for breast cancer regulate cyclin D1 expression through long-range enhancers. <i>American Journal of Human Genetics</i> , <b>2013</b> , 92, 489-503	11	167
156	Breast Cancer Risk Genes - Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , <b>2021</b> , 384, 428-439	59.2	143
155	Digital mammographic density and breast cancer risk: a case-control study of six alternative density assessment methods. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 439	8.3	141
154	CHEK2*1100delC heterozygosity in women with breast cancer associated with early death, breast cancer-specific death, and increased risk of a second breast cancer. <i>Journal of Clinical Oncology</i> , <b>2012</b> , 30, 4308-16	2.2	134
153	Genome-wide association study identifies a common variant associated with risk of endometrial cancer. <i>Nature Genetics</i> , <b>2011</b> , 43, 451-4	36.3	121
152	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , <b>2016</b> , 53, 800-811	5.8	121
151	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , <b>2018</b> , 50, 968-978	36.3	101

150	Mammographic density reduction is a prognostic marker of response to adjuvant tamoxifen therapy in postmenopausal patients with breast cancer. <i>Journal of Clinical Oncology</i> , <b>2013</b> , 31, 2249-56	2.2	96
149	Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. <i>Nature Genetics</i> , <b>2011</b> , 43, 185-7	36.3	96
148	Common breast cancer susceptibility variants in LSP1 and RAD51L1 are associated with mammographic density measures that predict breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 1156-66	4	92
147	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , <b>2014</b> , 4, 4999	17.4	87
146	A combined analysis of genome-wide association studies in breast cancer. <i>Breast Cancer Research and Treatment</i> , <b>2011</b> , 126, 717-27	4.4	85
145	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. <i>Nature Communications</i> , <b>2014</b> , 5, 5303	17.4	84
144	High-throughput mammographic-density measurement: a tool for risk prediction of breast cancer. <i>Breast Cancer Research</i> , <b>2012</b> , 14, R114	8.3	84
143	Risk factors and tumor characteristics of interval cancers by mammographic density. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, 1030-7	2.2	82
142	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> , <b>2013</b> , 93, 1046-60	11	80
141	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , <b>2016</b> , 13, e1002105	11.6	80
140	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	74
139	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , <b>2016</b> , 108,	9.7	65
138	Assessment of Breast Cancer Risk Factors Reveals Subtype Heterogeneity. <i>Cancer Research</i> , <b>2017</b> , 77, 3708-3717	10.1	64
137	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> , <b>2015</b> , 96, 5-20	11	59
136	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , <b>2016</b> , 48, 667-674	36.3	56
135	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , <b>2016</b> , 7, 12675	17.4	53
134	Automated measurement of volumetric mammographic density: a tool for widespread breast cancer risk assessment. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 1764-72	4	52
133	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , <b>2017</b> , 19, 599-603		51

132	Mammographic breast density and breast cancer: evidence of a shared genetic basis. <i>Cancer Research</i> , <b>2012</b> , 72, 1478-84	10.1	50
131	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , <b>2017</b> , 77, 2789-2799	10.1	49
130	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6096-111	5.6	48
129	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 1478-92	5.6	46
128	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , <b>2015</b> , 75, 2457-67	10.1	45
127	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. <i>American Journal of Human Genetics</i> , <b>2016</b> , 99, 903-911	11	43
126	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 1503-1510	4	42
125	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , <b>2016</b> , 23, 77-91	5.7	41
124	Change of mammographic density predicts the risk of contralateral breast cancer--a case-control study. <i>Breast Cancer Research</i> , <b>2013</b> , 15, R57	8.3	39
123	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25,000 subjects. <i>Molecular Psychiatry</i> , <b>2015</b> , 20, 735-43	15.1	39
122	Identification of novel genetic markers of breast cancer survival. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	38
121	Genetic predisposition to in situ and invasive lobular carcinoma of the breast. <i>PLoS Genetics</i> , <b>2014</b> , 10, e1004285	6	38
120	MicroRNA related polymorphisms and breast cancer risk. <i>PLoS ONE</i> , <b>2014</b> , 9, e109973	3.7	37
119	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 2966-84	5.6	36
118	Coffee consumption modifies risk of estrogen-receptor negative breast cancer. <i>Breast Cancer Research</i> , <b>2011</b> , 13, R49	8.3	36
117	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 285-98	5.6	35
116	Polygenic scores associated with educational attainment in adults predict educational achievement and ADHD symptoms in children. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2014</b> , 165B, 510-20	3.5	34
115	A genome-wide association scan on estrogen receptor-negative breast cancer. <i>Breast Cancer Research</i> , <b>2010</b> , 12, R93	8.3	32

114	Ethnic differences in mammographic densities: an Asian cross-sectional study. <i>PLoS ONE</i> , <b>2015</b> , 10, e0117568	3.5	31
113	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , <b>2020</b> , 11, 3833	17.4	31
112	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 22	8.3	31
111	Candidate locus analysis of the TERT-CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. <i>Human Genetics</i> , <b>2015</b> , 134, 231-45	6.3	30
110	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> , <b>2014</b> , 23, 1934-46	5.6	28
109	Identification of a novel percent mammographic density locus at 12q24. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 3299-305	5.6	28
108	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , <b>2021</b> , 596, 393-397	50.4	28
107	Associations of Breast Cancer Risk Prediction Tools With Tumor Characteristics and Metastasis. <i>Journal of Clinical Oncology</i> , <b>2016</b> , 34, 251-8	2.2	27
106	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , <b>2017</b> , 46, 1814-1822	7.8	27
105	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , <b>2015</b> , 5, 17369	4.9	27
104	The long-term prognostic and predictive capacity of cyclin D1 gene amplification in 2305 breast tumours. <i>Breast Cancer Research</i> , <b>2019</b> , 21, 34	8.3	26
103	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , <b>2015</b> , 97, 22-34	11	26
102	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , <b>2017</b> , 19, 119	8.3	26
101	A comprehensive tool for measuring mammographic density changes over time. <i>Breast Cancer Research and Treatment</i> , <b>2018</b> , 169, 371-379	4.4	26
100	Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. <i>International Journal of Cancer</i> , <b>2015</b> , 136, E685-96	7.5	26
99	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 1303-1317	7.5	26
98	Patient survival and tumor characteristics associated with CHEK2:p.I157T - findings from the Breast Cancer Association Consortium. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 98	8.3	26
97	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 64	8.3	25

96	Identification of new genetic susceptibility loci for breast cancer through consideration of gene-environment interactions. <i>Genetic Epidemiology</i> , <b>2014</b> , 38, 84-93	2.6	24
95	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 58	8.3	24
94	Breast cancer genetic risk profile is differentially associated with interval and screen-detected breast cancers. <i>Annals of Oncology</i> , <b>2015</b> , 26, 517-22	10.3	24
93	Area and volumetric density estimation in processed full-field digital mammograms for risk assessment of breast cancer. <i>PLoS ONE</i> , <b>2014</b> , 9, e110690	3.7	21
92	Common diseases as determinants of menopausal age. <i>Human Reproduction</i> , <b>2016</b> , 31, 2856-2864	5.7	20
91	FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. <i>British Journal of Cancer</i> , <b>2014</b> , 110, 1088-100	8.7	20
90	CYP2B6*6 is associated with increased breast cancer risk. <i>International Journal of Cancer</i> , <b>2014</b> , 134, 426-30	7.9	20
89	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. <i>Endocrine-Related Cancer</i> , <b>2015</b> , 22, 851-61	5.7	19
88	SNP-SNP interaction analysis of NF- $\kappa$ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , <b>2015</b> , 6, 37979-94	3.3	19
87	Blurring of high-resolution data shows that the effect of intrinsic nucleosome occupancy on transcription factor binding is mostly regional, not local. <i>PLoS Computational Biology</i> , <b>2010</b> , 6, e1000649 <sup>5</sup>	5	19
86	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> , <b>2019</b> , 21, 68	8.3	18
85	Identification of two novel mammographic density loci at 6Q25.1. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 75	8.3	18
84	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , <b>2016</b> , 11, e0153788	3.7	18
83	Worse quality of life in young and recently diagnosed breast cancer survivors compared with female survivors of other cancers: A cross-sectional study. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 2415-25	7.5	18
82	Prevalence of BRCA1 and BRCA2 pathogenic variants in a large, unselected breast cancer cohort. <i>International Journal of Cancer</i> , <b>2019</b> , 144, 1195-1204	7.5	18
81	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with prognosis of estrogen receptor-negative breast cancer after chemotherapy. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 18	8.3	17
80	Fine-scale mapping of the 4q24 locus identifies two independent loci associated with breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 1680-91	4	17
79	Impact of delayed treatment in women diagnosed with breast cancer: A population-based study. <i>Cancer Medicine</i> , <b>2020</b> , 9, 2435-2444	4.8	17

78	Mammographic Breast Density and Common Genetic Variants in Breast Cancer Risk Prediction. <i>PLoS ONE</i> , <b>2015</b> , 10, e0136650	3.7	17
77	9q31.2-rs865686 as a susceptibility locus for estrogen receptor-positive breast cancer: evidence from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 1783-91	4	17
76	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , <b>2020</b> , 11, 1217	17.4	16
75	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , <b>2019</b> , 48, 781-794	7.8	16
74	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , <b>2016</b> , 141, 386-401	4.9	15
73	Volumetric mammographic density: heritability and association with breast cancer susceptibility loci. <i>Journal of the National Cancer Institute</i> , <b>2014</b> , 106,	9.7	15
72	Enhancement of mammographic density measures in breast cancer risk prediction. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 1314-23	4	15
71	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , <b>2016</b> , 27, 679-93	2.8	15
70	A genome-wide association study to identify genetic susceptibility loci that modify ductal and lobular postmenopausal breast cancer risk associated with menopausal hormone therapy use: a two-stage design with replication. <i>Breast Cancer Research and Treatment</i> , <b>2013</b> , 138, 529-542	4.4	14
69	Genetic variation in the estrogen metabolic pathway and mammographic density as an intermediate phenotype of breast cancer. <i>Breast Cancer Research</i> , <b>2010</b> , 12, R19	8.3	14
68	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 110	8.3	13
67	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. <i>Nature Communications</i> , <b>2014</b> , 5, 4051	17.4	13
66	Differential Burden of Rare and Common Variants on Tumor Characteristics, Survival, and Mode of Detection in Breast Cancer. <i>Cancer Research</i> , <b>2018</b> , 78, 6329-6338	10.1	13
65	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> , <b>2015</b> , 36, 256-71	4.6	12
64	Molecular Differences between Screen-Detected and Interval Breast Cancers Are Largely Explained by PAM50 Subtypes. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 2584-2592	12.9	12
63	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , <b>2020</b> , 107, 837-848	11	12
62	The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. <i>European Journal of Epidemiology</i> , <b>2019</b> , 34, 591-600	12.1	11
61	Assessing within-woman changes in mammographic density: a comparison of fully versus semi-automated area-based approaches. <i>Cancer Causes and Control</i> , <b>2016</b> , 27, 481-91	2.8	11

60	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6034-46	5.6	11
59	Large-scale genotyping identifies a new locus at 22q13.2 associated with female breast size. <i>Journal of Medical Genetics</i> , <b>2013</b> , 50, 666-73	5.8	11
58	Calcium intake is not related to breast cancer risk among Singapore Chinese women. <i>International Journal of Cancer</i> , <b>2013</b> , 133, 680-6	7.5	11
57	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , <b>2016</b> , 11, e0160316	3.7	11
56	Impact of deviation from guideline recommended treatment on breast cancer survival in Asia. <i>Scientific Reports</i> , <b>2020</b> , 10, 1330	4.9	10
55	Determinants of breast size in Asian women. <i>Scientific Reports</i> , <b>2018</b> , 8, 1201	4.9	10
54	Common genetic variation and novel loci associated with volumetric mammographic density. <i>Breast Cancer Research</i> , <b>2018</b> , 20, 30	8.3	10
53	Assessment of a fully automated, high-throughput mammographic density measurement tool for use with processed digital mammograms. <i>Cancer Causes and Control</i> , <b>2014</b> , 25, 1037-43	2.8	10
52	The aetiology of convulsive status epilepticus: a study of 258 cases in Western China. <i>Seizure: the Journal of the British Epilepsy Association</i> , <b>2014</b> , 23, 717-21	3.2	10
51	Re-evaluating genetic variants identified in candidate gene studies of breast cancer risk using data from nearly 280,000 women of Asian and European ancestry. <i>EBioMedicine</i> , <b>2019</b> , 48, 203-211	8.8	9
50	Body size in early life and risk of breast cancer. <i>Breast Cancer Research</i> , <b>2017</b> , 19, 84	8.3	9
49	Inherited factors contribute to an inverse association between preeclampsia and breast cancer. <i>Breast Cancer Research</i> , <b>2018</b> , 20, 6	8.3	9
48	Confirmation of the reduction of hormone replacement therapy-related breast cancer risk for carriers of the HSD17B1_937_G variant. <i>Breast Cancer Research and Treatment</i> , <b>2013</b> , 138, 543-8	4.4	9
47	Association of CYP2D6 metabolizer status with mammographic density change in response to tamoxifen treatment. <i>Breast Cancer Research</i> , <b>2013</b> , 15, R93	8.3	9
46	Effects of childhood body size on breast cancer tumour characteristics. <i>Breast Cancer Research</i> , <b>2010</b> , 12, R23	8.3	9
45	Risk and predictors of psoriasis in patients with breast cancer: a Swedish population-based cohort study. <i>BMC Medicine</i> , <b>2017</b> , 15, 154	11.4	8
44	Treatment of Colon Cancer by Degradable rrPPC Nano-Conjugates Delivered STAT3 siRNA. <i>International Journal of Nanomedicine</i> , <b>2020</b> , 15, 9875-9890	7.3	8
43	Comparison of self-reported and register-based hospital medical data on comorbidities in women. <i>Scientific Reports</i> , <b>2019</b> , 9, 3527	4.9	7



42	The UGT1A6_19_GG genotype is a breast cancer risk factor. <i>Frontiers in Genetics</i> , <b>2013</b> , 4, 104	4.5	7
41	Factors associated with false-positive mammography at first screen in an Asian population. <i>PLoS ONE</i> , <b>2019</b> , 14, e0213615	3.7	6
40	Incidence of breast cancer attributable to breast density, modifiable and non-modifiable breast cancer risk factors in Singapore. <i>Scientific Reports</i> , <b>2020</b> , 10, 503	4.9	6
39	Variants in 6q25.1 Are Associated with Mammographic Density in Malaysian Chinese Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 327-33	4	6
38	Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. <i>Human Genetics</i> , <b>2016</b> , 135, 137-54	6.3	6
37	A polymorphism in the base excision repair gene PARP2 is associated with differential prognosis by chemotherapy among postmenopausal breast cancer patients. <i>BMC Cancer</i> , <b>2015</b> , 15, 978	4.8	6
36	The genetic basis of quality of life in healthy Swedish women: a candidate gene approach. <i>PLoS ONE</i> , <b>2015</b> , 10, e0118292	3.7	6
35	Functionalized DMP-039 Hybrid Nanoparticle as a Novel mRNA Vector for Efficient Cancer Suicide Gene Therapy. <i>International Journal of Nanomedicine</i> , <b>2021</b> , 16, 5211-5232	7.3	6
34	Family History, Reproductive, and Lifestyle Risk Factors for Fibroadenoma and Breast Cancer. <i>JNCI Cancer Spectrum</i> , <b>2018</b> , 2, pky051	4.6	5
33	The postmenopausal hormone replacement therapy-related breast cancer risk is decreased in women carrying the CYP2C19*17 variant. <i>Breast Cancer Research and Treatment</i> , <b>2012</b> , 131, 347-50	4.4	4
32	Longitudinal fluctuation in mammographic percent density differentiates between interval and screen-detected breast cancer. <i>International Journal of Cancer</i> , <b>2017</b> , 140, 34-40	7.5	4
31	Associations between childhood body size and seventeen adverse outcomes: analysis of 65,057 European women. <i>Scientific Reports</i> , <b>2017</b> , 7, 16917	4.9	4
30	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes.. <i>JAMA Oncology</i> , <b>2022</b> ,	13.4	4
29	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 623-642	4	4
28	Prognostic information of a previously diagnosed sister is an independent prognosticator for a newly diagnosed sister with breast cancer. <i>Annals of Oncology</i> , <b>2014</b> , 25, 1966-1972	10.3	3
27	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , <b>2019</b> , 9, 12524	4.9	2
26	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , <b>2020</b> , 10, 9688	4.9	2
25	rs2735383, located at a microRNA binding site in the 3'UTR of NBS1, is not associated with breast cancer risk. <i>Scientific Reports</i> , <b>2016</b> , 6, 36874	4.9	2

24	No evidence for association of inherited variation in genes involved in mitosis and percent mammographic density. <i>Breast Cancer Research</i> , <b>2012</b> , 14, R7	8.3	2
23	Common shared genetic variation behind decreased risk of breast cancer in celiac disease. <i>Scientific Reports</i> , <b>2017</b> , 7, 5942	4.9	2
22	DNA methylation and breast cancer-associated variants. <i>Breast Cancer Research and Treatment</i> , <b>2021</b> , 188, 713-727	4.4	2
21	Cohort profile: The Singapore Breast Cancer Cohort (SGBCC), a multi-center breast cancer cohort for evaluation of phenotypic risk factors and genetic markers. <i>PLoS ONE</i> , <b>2021</b> , 16, e0250102	3.7	2
20	Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. <i>Human Genetics and Genomics Advances</i> , <b>2021</b> , 2, 100042-100042	0.8	2
19	Epidemiological and ES cell-based functional evaluation of BRCA2 variants identified in families with breast cancer. <i>Human Mutation</i> , <b>2021</b> , 42, 200-212	4.7	2
18	Association between breast cancer risk and disease aggressiveness: Characterizing underlying gene expression patterns. <i>International Journal of Cancer</i> , <b>2021</b> , 148, 884-894	7.5	2
17	Polygenic risk scores for prediction of breast cancer risk in Asian populations.. <i>Genetics in Medicine</i> , <b>2021</b> ,	8.1	2
16	Automated breast tissue density assessment using high order regional texture descriptors in mammography <b>2014</b> ,		1
15	Predicting the Likelihood of Carrying a or Mutation in Asian Patients With Breast Cancer.. <i>Journal of Clinical Oncology</i> , <b>2022</b> , JCO2101647	2.2	1
14	In Search for the Genetic Basis of Quality of Life in Healthy Swedish Women--A GWAS Study Using the iCOGS Custom Genotyping Array. <i>PLoS ONE</i> , <b>2015</b> , 10, e0140563	3.7	1
13	Interval breast cancer is associated with interferon immune response.. <i>European Journal of Cancer</i> , <b>2022</b> , 162, 194-205	7.5	1
12	Breast cancer risk stratification for mammographic screening: A nation-wide screening cohort of 24,431 women in Singapore. <i>Cancer Medicine</i> , <b>2021</b> , 10, 8182-8191	4.8	1
11	Characterisation of protein-truncating and missense variants in in 15 768 women from Malaysia and Singapore. <i>Journal of Medical Genetics</i> , <b>2021</b> ,	5.8	1
10	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> , <b>2021</b> , 108, 1190-1203	11	1
9	Single Micelle Vectors based on Lipid/Block Copolymer Compositions as mRNA Formulations for Efficient Cancer Immunogene Therapy. <i>Molecular Pharmaceutics</i> , <b>2021</b> , 18, 4029-4045	5.6	1
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7	Germline breast cancer susceptibility genes, tumor characteristics, and survival. <i>Genome Medicine</i> , <b>2021</b> , 13, 185	14.4	0

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5	Overlap of high-risk individuals predicted by family history, and genetic and non-genetic breast cancer risk prediction models: implications for risk stratification.. <i>BMC Medicine</i> , <b>2022</b> , 20, 150	11.4	○
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