Jingmei Li

List of Publications by Year in Descending Order

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	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes.. *JAMA Oncology*, **2022**,	13.4	4
166	Predicting the Likelihood of Carrying a or Mutation in Asian Patients With Breast Cancer.. *Journal of Clinical Oncology*, **2022**, JCO2101647	2.2	1
165	Interval breast cancer is associated with interferon immune response.. *European Journal of Cancer*, **2022**, 162, 194-205	7.5	1
164	BREAst screening Tailored for HEr (BREATHE)-A study protocol on personalised risk-based breast cancer screening programme.. *PLoS ONE*, **2022**, 17, e0265965	3.7	1
163	Dual-RNA controlled delivery system inhibited tumor growth by apoptosis induction and TME activation.. *Journal of Controlled Release*, **2022**, 344, 97-112	11.7	0
162	Overlap of high-risk individuals predicted by family history, and genetic and non-genetic breast cancer risk prediction models: implications for risk stratification.. *BMC Medicine*, **2022**, 20, 150	11.4	0
161	Relevance of the MHC region for breast cancer susceptibility in Asians.. *Breast Cancer*, **2022**, 1	3.4	
160	Breast cancer risks associated with missense variants in breast cancer susceptibility genes.. *Genome Medicine*, **2022**, 14, 51	14.4	0
159	Association between Breast Cancer Polygenic Risk Score and Chemotherapy-Induced Febrile Neutropenia: Null Results. *Cancers*, **2022**, 14, 2714	6.6	0
158	Germline breast cancer susceptibility genes, tumor characteristics, and survival. *Genome Medicine*, **2021**, 13, 185	14.4	0
157	Breast cancer risk stratification for mammographic screening: A nation-wide screening cohort of 24,431 women in Singapore. *Cancer Medicine*, **2021**, 10, 8182-8191	4.8	1
156	DNA methylation and breast cancer-associated variants. *Breast Cancer Research and Treatment*, **2021**, 188, 713-727	4.4	2
155	Characterisation of protein-truncating and missense variants in in 15 768 women from Malaysia and Singapore. *Journal of Medical Genetics*, **2021**,	5.8	1
154	Cohort profile: The Singapore Breast Cancer Cohort (SGBCC), a multi-center breast cancer cohort for evaluation of phenotypic risk factors and genetic markers. *PLoS ONE*, **2021**, 16, e0250102	3.7	2
153	Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. *Human Genetics and Genomics Advances*, **2021**, 2, 100042-100042	0.8	2
152	Functionalized DMP-039 Hybrid Nanoparticle as a Novel mRNA Vector for Efficient Cancer Suicide Gene Therapy. *International Journal of Nanomedicine*, **2021**, 16, 5211-5232	7.3	6
151	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. *American Journal of Human Genetics*, **2021**, 108, 1190-1203	11	1

150	Epidemiological and ES cell-based functional evaluation of BRCA2 variants identified in families with breast cancer. *Human Mutation*, **2021**, 42, 200-212	4.7	2
149	Association between breast cancer risk and disease aggressiveness: Characterizing underlying gene expression patterns. *International Journal of Cancer*, **2021**, 148, 884-894	7.5	2
148	Breast Cancer Risk Genes - Association Analysis in More than 113,000 Women. *New England Journal of Medicine*, **2021**, 384, 428-439	59.2	143
147	Genetic insights into biological mechanisms governing human ovarian ageing. *Nature*, **2021**, 596, 393-397	50.4	28
146	Single Micelle Vectors based on Lipid/Block Copolymer Compositions as mRNA Formulations for Efficient Cancer Immunogene Therapy. *Molecular Pharmaceutics*, **2021**, 18, 4029-4045	5.6	1
145	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. *Cancer Epidemiology Biomarkers and Prevention*, **2021**, 30, 623-642	4	4
144	Polygenic risk scores for prediction of breast cancer risk in Asian populations.. *Genetics in Medicine*, **2021**,	8.1	2
143	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. *Scientific Reports*, **2020**, 10, 9688	4.9	2
142	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. *Nature Communications*, **2020**, 11, 1217	17.4	16
141	Impact of delayed treatment in women diagnosed with breast cancer: A population-based study. *Cancer Medicine*, **2020**, 9, 2435-2444	4.8	17
140	Incidence of breast cancer attributable to breast density, modifiable and non-modifiable breast cancer risk factors in Singapore. *Scientific Reports*, **2020**, 10, 503	4.9	6
139	Impact of deviation from guideline recommended treatment on breast cancer survival in Asia. *Scientific Reports*, **2020**, 10, 1330	4.9	10
138	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. *Nature Communications*, **2020**, 11, 3833	17.4	31
137	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. *American Journal of Human Genetics*, **2020**, 107, 837-848	11	12
136	Treatment of Colon Cancer by Degradable rrPPC Nano-Conjugates Delivered STAT3 siRNA. *International Journal of Nanomedicine*, **2020**, 15, 9875-9890	7.3	8
135	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. *EBioMedicine*, **2019**, 48, 203-211	8.8	9
134	Two truncating variants in FANCC and breast cancer risk. *Scientific Reports*, **2019**, 9, 12524	4.9	2
133	Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. *Breast Cancer Research*, **2019**, 21, 68	8.3	18

132	The long-term prognostic and predictive capacity of cyclin D1 gene amplification in 2305 breast tumours. *Breast Cancer Research*, **2019**, 21, 34	8.3	26
131	Comparison of self-reported and register-based hospital medical data on comorbidities in women. *Scientific Reports*, **2019**, 9, 3527	4.9	7
130	Factors associated with false-positive mammography at first screen in an Asian population. *PLoS ONE*, **2019**, 14, e0213615	3.7	6
129	The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. *European Journal of Epidemiology*, **2019**, 34, 591-600	12.1	11
128	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. *International Journal of Epidemiology*, **2019**, 48, 781-794	7.8	16
127	Prevalence of BRCA1 and BRCA2 pathogenic variants in a large, unselected breast cancer cohort. *International Journal of Cancer*, **2019**, 144, 1195-1204	7.5	18
126	A comprehensive tool for measuring mammographic density changes over time. *Breast Cancer Research and Treatment*, **2018**, 169, 371-379	4.4	26
125	Determinants of breast size in Asian women. *Scientific Reports*, **2018**, 8, 1201	4.9	10
124	Inherited factors contribute to an inverse association between preeclampsia and breast cancer. *Breast Cancer Research*, **2018**, 20, 6	8.3	9
123	Common genetic variation and novel loci associated with volumetric mammographic density. *Breast Cancer Research*, **2018**, 20, 30	8.3	10
122	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. *Nature Genetics*, **2018**, 50, 968-978	36.3	101
121	Family History, Reproductive, and Lifestyle Risk Factors for Fibroadenoma and Breast Cancer. *JNCI Cancer Spectrum*, **2018**, 2, pky051	4.6	5
120	Differential Burden of Rare and Common Variants on Tumor Characteristics, Survival, and Mode of Detection in Breast Cancer. *Cancer Research*, **2018**, 78, 6329-6338	10.1	13
119	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. *Cancer Research*, **2017**, 77, 2789-2799	10.1	49
118	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. *Nature Genetics*, **2017**, 49, 834-841	36.3	257
117	Assessment of Breast Cancer Risk Factors Reveals Subtype Heterogeneity. *Cancer Research*, **2017**, 77, 3708-3717	10.1	64
116	Association analysis identifies 65 new breast cancer risk loci. *Nature*, **2017**, 551, 92-94	50.4	643
115	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. *Nature Genetics*, **2017**, 49, 1767-1778	36.3	186

114	Risk and predictors of psoriasis in patients with breast cancer: a Swedish population-based cohort study. *BMC Medicine*, **2017**, 15, 154	11.4	8
113	Body size in early life and risk of breast cancer. *Breast Cancer Research*, **2017**, 19, 84	8.3	9
112	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. *Breast Cancer Research*, **2017**, 19, 119	8.3	26
111	Common shared genetic variation behind decreased risk of breast cancer in celiac disease. *Scientific Reports*, **2017**, 7, 5942	4.9	2
110	Longitudinal fluctuation in mammographic percent density differentiates between interval and screen-detected breast cancer. *International Journal of Cancer*, **2017**, 140, 34-40	7.5	4
109	Molecular Differences between Screen-Detected and Interval Breast Cancers Are Largely Explained by PAM50 Subtypes. *Clinical Cancer Research*, **2017**, 23, 2584-2592	12.9	12
108	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. *Genetics in Medicine*, **2017**, 19, 599-603	6.3	51
107	Associations between childhood body size and seventeen adverse outcomes: analysis of 65,057 European women. *Scientific Reports*, **2017**, 7, 16917	4.9	4
106	Body mass index and breast cancer survival: a Mendelian randomization analysis. *International Journal of Epidemiology*, **2017**, 46, 1814-1822	7.8	27
105	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. *Gynecologic Oncology*, **2016**, 141, 386-401	4.9	15
104	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2016**, 25, 1503-1510	4	42
103	rs2735383, located at a microRNA binding site in the 3′UTR of NBS1, is not associated with breast cancer risk. *Scientific Reports*, **2016**, 6, 36874	4.9	2
102	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. *Nature Communications*, **2016**, 7, 12675	17.4	53
101	Common diseases as determinants of menopausal age. *Human Reproduction*, **2016**, 31, 2856-2864	5.7	20
100	Assessing within-woman changes in mammographic density: a comparison of fully versus semi-automated area-based approaches. *Cancer Causes and Control*, **2016**, 27, 481-91	2.8	11
99	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. *Endocrine-Related Cancer*, **2016**, 23, 77-91	5.7	41
98	Variants in 6q25.1 Are Associated with Mammographic Density in Malaysian Chinese Women. *Cancer Epidemiology Biomarkers and Prevention*, **2016**, 25, 327-33	4	6
97	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. *Human Genetics*, **2016**, 135, 137-54	6.3	6

96	Associations of Breast Cancer Risk Prediction Tools With Tumor Characteristics and Metastasis. *Journal of Clinical Oncology*, **2016**, 34, 251-8	2.2	27
95	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. *Journal of the National Cancer Institute*, **2016**, 108,	9.7	65
94	RAD51B in Familial Breast Cancer. *PLoS ONE*, **2016**, 11, e0153788	3.7	18
93	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. *PLoS Medicine*, **2016**, 13, e1002105	11.6	80
92	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. *PLoS ONE*, **2016**, 11, e0160316	3.7	11
91	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. *International Journal of Cancer*, **2016**, 139, 1303-1317	7.5	26
90	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. *Journal of Medical Genetics*, **2016**, 53, 800-811	5.8	121
89	Patient survival and tumor characteristics associated with CHEK2:p.I157T - findings from the Breast Cancer Association Consortium. *Breast Cancer Research*, **2016**, 18, 98	8.3	26
88	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. *Breast Cancer Research*, **2016**, 18, 64	8.3	25
87	Genetic predisposition to ductal carcinoma in situ of the breast. *Breast Cancer Research*, **2016**, 18, 22	8.3	31
86	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. *Cancer Causes and Control*, **2016**, 27, 679-93	2.8	15
85	Five endometrial cancer risk loci identified through genome-wide association analysis. *Nature Genetics*, **2016**, 48, 667-674	36.3	56
84	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. *American Journal of Human Genetics*, **2016**, 99, 903-911	11	43
83	Worse quality of life in young and recently diagnosed breast cancer survivors compared with female survivors of other cancers: A cross-sectional study. *International Journal of Cancer*, **2016**, 139, 2415-25	7.5	18
82	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. *Carcinogenesis*, **2015**, 36, 256-71	4.6	12
81	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. *Nature Genetics*, **2015**, 47, 373-80	36.3	406
80	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. *American Journal of Human Genetics*, **2015**, 97, 22-34	11	26
79	Identification of novel genetic markers of breast cancer survival. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	38

78	Candidate locus analysis of the TERT-CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. *Human Genetics*, **2015**, 134, 231-45	6.3	30
77	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with prognosis of estrogen receptor-negative breast cancer after chemotherapy. *Breast Cancer Research*, **2015**, 17, 18	8.3	17
76	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. *Cancer Research*, **2015**, 75, 2457-67	10.1	45
75	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. *Nature Genetics*, **2015**, 47, 1294-1303	36.3	226
74	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	74
73	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. *Endocrine-Related Cancer*, **2015**, 22, 851-61	5.7	19
72	Fine-scale mapping of the 4q24 locus identifies two independent loci associated with breast cancer risk. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 1680-91	4	17
71	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. *Human Molecular Genetics*, **2015**, 24, 285-98	5.6	35
70	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. *Human Molecular Genetics*, **2015**, 24, 1478-92	5.6	46
69	Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. *International Journal of Cancer*, **2015**, 136, E685-96	7.5	26
68	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. *Scientific Reports*, **2015**, 5, 17369	4.9	27
67	A polymorphism in the base excision repair gene PARP2 is associated with differential prognosis by chemotherapy among postmenopausal breast cancer patients. *BMC Cancer*, **2015**, 15, 978	4.8	6
66	Common germline polymorphisms associated with breast cancer-specific survival. *Breast Cancer Research*, **2015**, 17, 58	8.3	24
65	Identification of two novel mammographic density loci at 6Q25.1. *Breast Cancer Research*, **2015**, 17, 75	8.3	18
64	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. *Breast Cancer Research*, **2015**, 17, 110	8.3	13
63	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. *Obstetrical and Gynecological Survey*, **2015**, 70, 758-762	2.4	
62	The genetic basis of quality of life in healthy Swedish women: a candidate gene approach. *PLoS ONE*, **2015**, 10, e0118292	3.7	6
61	Mammographic Breast Density and Common Genetic Variants in Breast Cancer Risk Prediction. *PLoS ONE*, **2015**, 10, e0136650	3.7	17

60	SNP-SNP interaction analysis of NF-B signaling pathway on breast cancer survival. *Oncotarget*, **2015**, 6, 37979-94	3.3	19
59	Prediction of breast cancer risk based on profiling with common genetic variants. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	324
58	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25,000 subjects. *Molecular Psychiatry*, **2015**, 20, 735-43	15.1	39
57	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. *Human Molecular Genetics*, **2015**, 24, 2966-84	5.6	36
56	Risk factors and tumor characteristics of interval cancers by mammographic density. *Journal of Clinical Oncology*, **2015**, 33, 1030-7	2.2	82
55	Fine-scale mapping of the 5q11.2 breast cancer locus reveals at least three independent risk variants regulating MAP3K1. *American Journal of Human Genetics*, **2015**, 96, 5-20	11	59
54	Breast cancer genetic risk profile is differentially associated with interval and screen-detected breast cancers. *Annals of Oncology*, **2015**, 26, 517-22	10.3	24
53	Ethnic differences in mammographic densities: an Asian cross-sectional study. *PLoS ONE*, **2015**, 10, e0117568	3.7	31
52	In Search for the Genetic Basis of Quality of Life in Healthy Swedish Women--A GWAS Study Using the iCOGS Custom Genotyping Array. *PLoS ONE*, **2015**, 10, e0140563	3.7	1
51	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. *Human Molecular Genetics*, **2014**, 23, 1934-46	5.6	28
50	Identification of new genetic susceptibility loci for breast cancer through consideration of gene-environment interactions. *Genetic Epidemiology*, **2014**, 38, 84-93	2.6	24
49	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. *Nature Communications*, **2014**, 5, 5303	17.4	84
48	Mammographic density phenotypes and risk of breast cancer: a meta-analysis. *Journal of the National Cancer Institute*, **2014**, 106,	9.7	190
47	FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. *British Journal of Cancer*, **2014**, 110, 1088-100	8.7	20
46	Assessment of a fully automated, high-throughput mammographic density measurement tool for use with processed digital mammograms. *Cancer Causes and Control*, **2014**, 25, 1037-43	2.8	10
45	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. *Nature Communications*, **2014**, 4, 4999	17.4	87
44	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. *Human Molecular Genetics*, **2014**, 23, 6034-46	5.6	11
43	The aetiology of convulsive status epilepticus: a study of 258 cases in Western China. *Seizure: the Journal of the British Epilepsy Association*, **2014**, 23, 717-21	3.2	10

42	MicroRNA related polymorphisms and breast cancer risk. *PLoS ONE*, **2014**, 9, e109973	3.7	37
41	Area and volumetric density estimation in processed full-field digital mammograms for risk assessment of breast cancer. *PLoS ONE*, **2014**, 9, e110690	3.7	21
40	Automated measurement of volumetric mammographic density: a tool for widespread breast cancer risk assessment. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 1764-72	4	52
39	Digital mammographic density and breast cancer risk: a case-control study of six alternative density assessment methods. *Breast Cancer Research*, **2014**, 16, 439	8.3	141
38	Volumetric mammographic density: heritability and association with breast cancer susceptibility loci. *Journal of the National Cancer Institute*, **2014**, 106,	9.7	15
37	Genetic predisposition to in situ and invasive lobular carcinoma of the breast. *PLoS Genetics*, **2014**, 10, e1004285	6	38
36	Enhancement of mammographic density measures in breast cancer risk prediction. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 1314-23	4	15
35	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. *Nature Communications*, **2014**, 5, 4051	17.4	13
34	Polygenic scores associated with educational attainment in adults predict educational achievement and ADHD symptoms in children. *American Journal of Medical Genetics Part B: Neuropsychiatric Genetics*, **2014**, 165B, 510-20	3.5	34
33	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. *Human Molecular Genetics*, **2014**, 23, 6096-111	5.6	48
32	Prognostic information of a previously diagnosed sister is an independent prognosticator for a newly diagnosed sister with breast cancer. *Annals of Oncology*, **2014**, 25, 1966-1972	10.3	3
31	Automated breast tissue density assessment using high order regional texture descriptors in mammography **2014**,		1
30	CYP2B6*6 is associated with increased breast cancer risk. *International Journal of Cancer*, **2014**, 134, 426-39	7.9	20
29	Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1. *American Journal of Human Genetics*, **2013**, 93, 1046-60	11	80
28	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. *Nature Genetics*, **2013**, 45, 371-84, 384e1-2	36.3	422
27	Change of mammographic density predicts the risk of contralateral breast cancer--a case-control study. *Breast Cancer Research*, **2013**, 15, R57	8.3	39
26	Confirmation of the reduction of hormone replacement therapy-related breast cancer risk for carriers of the HSD17B1_937_G variant. *Breast Cancer Research and Treatment*, **2013**, 138, 543-8	4.4	9
25	Functional variants at the 11q13 risk locus for breast cancer regulate cyclin D1 expression through long-range enhancers. *American Journal of Human Genetics*, **2013**, 92, 489-503	11	167

24	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. *Nature Genetics*, **2013**, 45, 353-61, 361e1-2	36.3	813
23	GWAS of 126,559 individuals identifies genetic variants associated with educational attainment. *Science*, **2013**, 340, 1467-71	33.3	563
22	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. *Breast Cancer Research and Treatment*, **2013**, 138, 529-542	4.4	14
21	Mammographic density reduction is a prognostic marker of response to adjuvant tamoxifen therapy in postmenopausal patients with breast cancer. *Journal of Clinical Oncology*, **2013**, 31, 2249-56	2.2	96
20	Large-scale genotyping identifies a new locus at 22q13.2 associated with female breast size. *Journal of Medical Genetics*, **2013**, 50, 666-73	5.8	11
19	Calcium intake is not related to breast cancer risk among Singapore Chinese women. *International Journal of Cancer*, **2013**, 133, 680-6	7.5	11
18	Association of CYP2D6 metabolizer status with mammographic density change in response to tamoxifen treatment. *Breast Cancer Research*, **2013**, 15, R93	8.3	9
17	The UGT1A6_19_GG genotype is a breast cancer risk factor. *Frontiers in Genetics*, **2013**, 4, 104	4.5	7
16	No evidence for association of inherited variation in genes involved in mitosis and percent mammographic density. *Breast Cancer Research*, **2012**, 14, R7	8.3	2
15	The postmenopausal hormone replacement therapy-related breast cancer risk is decreased in women carrying the CYP2C19*17 variant. *Breast Cancer Research and Treatment*, **2012**, 131, 347-50	4.4	4
14	High-throughput mammographic-density measurement: a tool for risk prediction of breast cancer. *Breast Cancer Research*, **2012**, 14, R114	8.3	84
13	9q31.2-rs865686 as a susceptibility locus for estrogen receptor-positive breast cancer: evidence from the Breast Cancer Association Consortium. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 1783-91	4	17
12	Common breast cancer susceptibility variants in LSP1 and RAD51L1 are associated with mammographic density measures that predict breast cancer risk. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 1156-66	4	92
11	Identification of a novel percent mammographic density locus at 12q24. *Human Molecular Genetics*, **2012**, 21, 3299-305	5.6	28
10	Mammographic breast density and breast cancer: evidence of a shared genetic basis. *Cancer Research*, **2012**, 72, 1478-84	10.1	50
9	CHEK2*1100delC heterozygosity in women with breast cancer associated with early death, breast cancer-specific death, and increased risk of a second breast cancer. *Journal of Clinical Oncology*, **2012**, 30, 4308-16	2.2	134
8	Coffee consumption modifies risk of estrogen-receptor negative breast cancer. *Breast Cancer Research*, **2011**, 13, R49	8.3	36
7	Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. *Nature Genetics*, **2011**, 43, 185-7	36.3	96

6	Genome-wide association study identifies a common variant associated with risk of endometrial cancer. *Nature Genetics*, **2011**, 43, 451-4	36.3	121
5	A combined analysis of genome-wide association studies in breast cancer. *Breast Cancer Research and Treatment*, **2011**, 126, 717-27	4.4	85
4	Blurring of high-resolution data shows that the effect of intrinsic nucleosome occupancy on transcription factor binding is mostly regional, not local. *PLoS Computational Biology*, **2010**, 6, e1000649	5	19
3	A genome-wide association scan on estrogen receptor-negative breast cancer. *Breast Cancer Research*, **2010**, 12, R93	8.3	32
2	Effects of childhood body size on breast cancer tumour characteristics. *Breast Cancer Research*, **2010**, 12, R23	8.3	9
1	Genetic variation in the estrogen metabolic pathway and mammographic density as an intermediate phenotype of breast cancer. *Breast Cancer Research*, **2010**, 12, R19	8.3	14