

# Jingmei Li

## List of Publications by Year in descending order

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Version: 2024-02-01

172  
papers

12,182  
citations

44069

48  
h-index

32842

100  
g-index

181  
all docs

181  
docs citations

181  
times ranked

17329  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.   | 27.8 | 1,099     |
| 2  | Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.   | 21.4 | 960       |
| 3  | GWAS of 126,559 Individuals Identifies Genetic Variants Associated with Educational Attainment. <i>Science</i> , 2013, 340, 1467-1471.  | 12.6 | 750       |
| 4  | Breast Cancer Risk Genes " Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.  | 27.0 | 532       |
| 5  | Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.  | 21.4 | 513       |
| 6  | Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.   | 21.4 | 493       |
| 7  | Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .  | 6.3  | 428       |
| 8  | 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> , 2017, 49, 834-841.  | 21.4 | 426       |
| 9  | Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.   | 21.4 | 357       |
| 10 | Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.  | 21.4 | 289       |
| 11 | Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. <i>Journal of the National Cancer Institute</i> , 2014, 106, .  | 6.3  | 261       |
| 12 | 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> , 2013, 92, 489-503.  | 6.2  | 201       |
| 13 | A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.  | 21.4 | 184       |
| 14 | Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.   | 27.8 | 183       |
| 15 | <i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.   | 3.2  | 174       |
| 16 | Digital mammographic density and breast cancer risk: a case-control study of six alternative density assessment methods. <i>Breast Cancer Research</i> , 2014, 16, 439.   | 5.0  | 165       |
| 17 | <i>CHEK2</i> *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> , 2012, 30, 4308-4316. | 1.6  | 162       |
| 18 | Genome-wide association study identifies a common variant associated with risk of endometrial cancer. <i>Nature Genetics</i> , 2011, 43, 451-454.   | 21.4 | 141       |

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|----|---|------|-----------|
| 19 | 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> , 2016, 13, e1002105.   | 8.4  | 118       |
| 20 | 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> , 2013, 31, 2249-2256.                                     | 1.6  | 113       |
| 21 | Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. <i>Nature Genetics</i> , 2011, 43, 185-187.   | 21.4 | 109       |
| 22 | Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. <i>Nature Communications</i> , 2014, 5, 5303.  | 12.8 | 109       |
| 23 | Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.   | 12.8 | 105       |
| 24 | Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166. | 2.5  | 101       |
| 25 | Risk Factors and Tumor Characteristics of Interval Cancers by Mammographic Density. <i>Journal of Clinical Oncology</i> , 2015, 33, 1030-1037.  | 1.6  | 99        |
| 26 | Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv219.  | 6.3  | 99        |
| 27 | Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.   | 6.2  | 98        |
| 28 | High-throughput mammographic-density measurement: a tool for risk prediction of breast cancer. <i>Breast Cancer Research</i> , 2012, 14, R114.  | 5.0  | 96        |
| 29 | A combined analysis of genome-wide association studies in breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011, 126, 717-727.  | 2.5  | 90        |
| 30 | European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833.  | 12.8 | 88        |
| 31 | Assessment of Breast Cancer Risk Factors Reveals Subtype Heterogeneity. <i>Cancer Research</i> , 2017, 77, 3708-3717.   | 0.9  | 87        |
| 32 | Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.   | 12.8 | 78        |
| 33 | Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016, 48, 667-674.  | 21.4 | 77        |
| 34 | BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv315.   | 6.3  | 77        |
| 35 | Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. <i>American Journal of Human Genetics</i> , 2015, 96, 5-20.  | 6.2  | 76        |
| 36 | <i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.  | 0.9  | 75        |

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|----|---|------|-----------|
| 37 | Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.   | 2.4  | 67        |
| 38 | 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> , 2016, 25, 1503-1510. | 2.5  | 64        |
| 39 | Automated Measurement of Volumetric Mammographic Density: A Tool for Widespread Breast Cancer Risk Assessment. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1764-1772.  | 2.5  | 62        |
| 40 | CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.  | 3.1  | 62        |
| 41 | The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25â€‰%000 subjects. <i>Molecular Psychiatry</i> , 2015, 20, 735-743.   | 7.9  | 59        |
| 42 | 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> , 2016, 99, 903-911.                  | 6.2  | 59        |
| 43 | Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .   | 6.3  | 56        |
| 44 | Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , 2015, 75, 2457-2467.  | 0.9  | 55        |
| 45 | Mammographic Breast Density and Breast Cancer: Evidence of a Shared Genetic Basis. <i>Cancer Research</i> , 2012, 72, 1478-1484.  | 0.9  | 54        |
| 46 | Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.   | 2.9  | 53        |
| 47 | Fineâ€‰scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.  | 5.1  | 51        |
| 48 | Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744.  | 7.1  | 51        |
| 49 | Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 1478-1492.  | 2.9  | 50        |
| 50 | MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.  | 2.5  | 49        |
| 51 | The long-term prognostic and predictive capacity of cyclin D1 gene amplification in 2305 breast tumours. <i>Breast Cancer Research</i> , 2019, 21, 34.  | 5.0  | 48        |
| 52 | Coffee consumption modifies risk of estrogen-receptor negative breast cancer. <i>Breast Cancer Research</i> , 2011, 13, R49.  | 5.0  | 46        |
| 53 | Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.   | 12.8 | 46        |
| 54 | Impact of delayed treatment in women diagnosed with breast cancer: A populationâ€‰based study. <i>Cancer Medicine</i> , 2020, 9, 2435-2444.   | 2.8  | 46        |

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|----|--|-----|-----------|
| 55 | Associations of Breast Cancer Risk Prediction Tools With Tumor Characteristics and Metastasis. <i>Journal of Clinical Oncology</i> , 2016, 34, 251-258.  | 1.6 | 45        |
| 56 | Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.  | 1.9 | 45        |
| 57 | A comprehensive tool for measuring mammographic density changes over time. <i>Breast Cancer Research and Treatment</i> , 2018, 169, 371-379.   | 2.5 | 45        |
| 58 | Change of mammographic density predicts the risk of contralateral breast cancer - a case-control study. <i>Breast Cancer Research</i> , 2013, 15, R57.   | 5.0 | 44        |
| 59 | Ethnic Differences in Mammographic Densities: An Asian Cross-Sectional Study. <i>PLoS ONE</i> , 2015, 10, e0117568.  | 2.5 | 44        |
| 60 | Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016, 18, 22.  | 5.0 | 43        |
| 61 | Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017, 19, 119.   | 5.0 | 43        |
| 62 | Common diseases as determinants of menopausal age. <i>Human Reproduction</i> , 2016, 31, 2856-2864.  | 0.9 | 42        |
| 63 | 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> , 2014, 165, 510-520. | 1.7 | 40        |
| 64 | Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.   | 2.9 | 40        |
| 65 | Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.   | 3.5 | 39        |
| 66 | Patient survival and tumor characteristics associated with CHEK2:p.I157T findings from the Breast Cancer Association Consortium. <i>Breast Cancer Research</i> , 2016, 18, 98.   | 5.0 | 39        |
| 67 | Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.   | 6.2 | 39        |
| 68 | Breast cancer genetic risk profile is differentially associated with interval and screen-detected breast cancers. <i>Annals of Oncology</i> , 2015, 26, 517-522.   | 1.2 | 38        |
| 69 | Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 285-298.   | 2.9 | 38        |
| 70 | Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , 2015, 97, 22-34.   | 6.2 | 37        |
| 71 | The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 781-794.   | 1.9 | 37        |
| 72 | A genome-wide association scan on estrogen receptor-negative breast cancer. <i>Breast Cancer Research</i> , 2010, 12, R93.   | 5.0 | 35        |

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|----|---|-----|-----------|
| 73 | Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369.                           | 3.3 | 35        |
| 74 | Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. <i>International Journal of Cancer</i> , 2015, 136, E685-96.                         | 5.1 | 34        |
| 75 | 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> , 2015, 134, 231-245.                  | 3.8 | 34        |
| 76 | A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46 450 cases and 42 461 controls from the breast cancer association consortium. <i>Human Molecular Genetics</i> , 2014, 23, 1934-1946. | 2.9 | 32        |
| 77 | Identification of a novel percent mammographic density locus at 12q24. <i>Human Molecular Genetics</i> , 2012, 21, 3299-3305.   | 2.9 | 31        |
| 78 | 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> , 2016, 18, 64.                            | 5.0 | 31        |
| 79 | Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. <i>Breast Cancer Research</i> , 2019, 21, 68.  | 5.0 | 31        |
| 80 | Prevalence of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants in a large, unselected breast cancer cohort. <i>International Journal of Cancer</i> , 2019, 144, 1195-1204.   | 5.1 | 31        |
| 81 | Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Gene-Environment Interactions. <i>Genetic Epidemiology</i> , 2014, 38, 84-93.  | 1.3 | 28        |
| 82 | 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> , 2010, 6, e1000649.                    | 3.2 | 27        |
| 83 | Polygenic risk scores for prediction of breast cancer risk in Asian populations. <i>Genetics in Medicine</i> , 2022, 24, 586-600.   | 2.4 | 27        |
| 84 | Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.  | 5.0 | 26        |
| 85 | RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.   | 2.5 | 26        |
| 86 | Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2015, 22, 851-861.  | 3.1 | 25        |
| 87 | Area and Volumetric Density Estimation in Processed Full-Field Digital Mammograms for Risk Assessment of Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e110690.   | 2.5 | 24        |
| 88 | <i>CYP2B6</i> *6 is associated with increased breast cancer risk. <i>International Journal of Cancer</i> , 2014, 134, 426-430.  | 5.1 | 24        |
| 89 | Identification of two novel mammographic density loci at 6Q25.1. <i>Breast Cancer Research</i> , 2015, 17, 75.  | 5.0 | 24        |
| 90 | Fine-Scale Mapping of the 4q24 Locus Identifies Two Independent Loci Associated with Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1680-1691.  | 2.5 | 24        |

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|-----|---|-----|-----------|
| 91  | Functionalized DMP-039 Hybrid Nanoparticle as a Novel mRNA Vector for Efficient Cancer Suicide Gene Therapy. <i>International Journal of Nanomedicine</i> , 2021, Volume 16, 5211-5232.   | 6.7 | 24        |
| 92  | 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> , 2016, 139, 2415-2425.  | 5.1 | 23        |
| 93  | Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.  | 2.8 | 23        |
| 94  | Treatment of Colon Cancer by Degradable rrPPC Nano-Conjugates Delivered STAT3 siRNA. <i>International Journal of Nanomedicine</i> , 2020, Volume 15, 9875-9890.   | 6.7 | 22        |
| 95  | Volumetric Mammographic Density: Heritability and Association With Breast Cancer Susceptibility Loci. <i>Journal of the National Cancer Institute</i> , 2014, 106, dju334-dju334.   | 6.3 | 21        |
| 96  | FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. <i>British Journal of Cancer</i> , 2014, 110, 1088-1100.  | 6.4 | 21        |
| 97  | Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016, 27, 679-693.   | 1.8 | 21        |
| 98  | Mammographic Breast Density and Common Genetic Variants in Breast Cancer Risk Prediction. <i>PLoS ONE</i> , 2015, 10, e0136650.   | 2.5 | 20        |
| 99  | SNP-SNP interaction analysis of NF- $\kappa$ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , 2015, 6, 37979-37994.   | 1.8 | 20        |
| 100 | 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> , 2015, 17, 18.   | 5.0 | 20        |
| 101 | A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. <i>Breast Cancer Research</i> , 2015, 17, 110.  | 5.0 | 19        |
| 102 | Determinants of breast size in Asian women. <i>Scientific Reports</i> , 2018, 8, 1201.  | 3.3 | 19        |
| 103 | Family History, Reproductive, and Lifestyle Risk Factors for Fibroadenoma and Breast Cancer. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky051.   | 2.9 | 19        |
| 104 | Differential Burden of Rare and Common Variants on Tumor Characteristics, Survival, and Mode of Detection in Breast Cancer. <i>Cancer Research</i> , 2018, 78, 6329-6338.   | 0.9 | 19        |
| 105 | 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.  | 2.5 | 19        |
| 106 | Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51.   | 8.2 | 19        |
| 107 | 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> , 2013, 138, 529-542. | 2.5 | 18        |
| 108 | No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.  | 1.4 | 18        |

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|-----|---|------|-----------|
| 109 | Common genetic variation and novel loci associated with volumetric mammographic density. <i>Breast Cancer Research</i> , 2018, 20, 30.  | 5.0  | 18        |
| 110 | 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> , 2012, 21, 1783-1791. | 2.5  | 17        |
| 111 | Enhancement of Mammographic Density Measures in Breast Cancer Risk Prediction. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1314-1323.  | 2.5  | 17        |
| 112 | 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.  | 5.0  | 16        |
| 113 | 2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. <i>Nature Communications</i> , 2014, 5, 4051.   | 12.8 | 16        |
| 114 | The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. <i>European Journal of Epidemiology</i> , 2019, 34, 591-600.  | 5.7  | 16        |
| 115 | Calcium intake is not related to breast cancer risk among Singapore Chinese women. <i>International Journal of Cancer</i> , 2013, 133, 680-686.   | 5.1  | 15        |
| 116 | Assessing within-woman changes in mammographic density: a comparison of fully versus semi-automated area-based approaches. <i>Cancer Causes and Control</i> , 2016, 27, 481-491.  | 1.8  | 15        |
| 117 | Molecular Differences between Screen-Detected and Interval Breast Cancers Are Largely Explained by PAM50 Subtypes. <i>Clinical Cancer Research</i> , 2017, 23, 2584-2592.   | 7.0  | 15        |
| 118 | Body size in early life and risk of breast cancer. <i>Breast Cancer Research</i> , 2017, 19, 84.  | 5.0  | 15        |
| 119 | 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.             | 2.8  | 14        |
| 120 | Inherited factors contribute to an inverse association between preeclampsia and breast cancer. <i>Breast Cancer Research</i> , 2018, 20, 6.   | 5.0  | 14        |
| 121 | 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> , 2019, 48, 203-211.                     | 6.1  | 14        |
| 122 | Incidence of breast cancer attributable to breast density, modifiable and non-modifiable breast cancer risk factors in Singapore. <i>Scientific Reports</i> , 2020, 10, 503.  | 3.3  | 14        |
| 123 | Impact of deviation from guideline recommended treatment on breast cancer survival in Asia. <i>Scientific Reports</i> , 2020, 10, 1330.   | 3.3  | 14        |
| 124 | Predicting the Likelihood of Carrying a <i>BRCA1</i> or <i>BRCA2</i> Mutation in Asian Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2022, 40, 1542-1551.  | 1.6  | 14        |
| 125 | Effects of childhood body size on breast cancer tumour characteristics. <i>Breast Cancer Research</i> , 2010, 12, R23.  | 5.0  | 13        |
| 126 | Risk and predictors of psoriasis in patients with breast cancer: a Swedish population-based cohort study. <i>BMC Medicine</i> , 2017, 15, 154.  | 5.5  | 13        |



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|-----|---|-----|-----------|
| 127 | Comparison of self-reported and register-based hospital medical data on comorbidities in women. <i>Scientific Reports</i> , 2019, 9, 3527.  | 3.3 | 13        |
| 128 | Single Micelle Vectors based on Lipid/Block Copolymer Compositions as mRNA Formulations for Efficient Cancer Immunogene Therapy. <i>Molecular Pharmaceutics</i> , 2021, 18, 4029-4045.  | 4.6 | 13        |
| 129 | Large-scale genotyping identifies a new locus at 22q13.2 associated with female breast size. <i>Journal of Medical Genetics</i> , 2013, 50, 666-673.  | 3.2 | 12        |
| 130 | Assessment of a fully automated, high-throughput mammographic density measurement tool for use with processed digital mammograms. <i>Cancer Causes and Control</i> , 2014, 25, 1037-1043.   | 1.8 | 12        |
| 131 | Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014, 23, 6034-6046.   | 2.9 | 12        |
| 132 | The aetiology of convulsive status epilepticus: A study of 258 cases in Western China. <i>Seizure: the Journal of the British Epilepsy Association</i> , 2014, 23, 717-721.   | 2.0 | 12        |
| 133 | Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.  | 2.5 | 12        |
| 134 | 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> , 2021, 16, e0250102.  | 2.5 | 11        |
| 135 | BREAst screening Tailored for HER (BREATHE) – A study protocol on personalised risk-based breast cancer screening programme. <i>PLoS ONE</i> , 2022, 17, e0265965.  | 2.5 | 11        |
| 136 | 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> , 2013, 138, 543-548.  | 2.5 | 10        |
| 137 | Association of CYP2D6 metabolizer status with mammographic density change in response to tamoxifen treatment. <i>Breast Cancer Research</i> , 2013, 15, R93.  | 5.0 | 10        |
| 138 | Variants in 6q25.1 Are Associated with Mammographic Density in Malaysian Chinese Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 327-333.   | 2.5 | 10        |
| 139 | Factors associated with false-positive mammography at first screen in an Asian population. <i>PLoS ONE</i> , 2019, 14, e0213615.  | 2.5 | 9         |
| 140 | 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> , 2022, 20, 150.                                  | 5.5 | 9         |
| 141 | The UGT1A6_19_GG genotype is a breast cancer risk factor. <i>Frontiers in Genetics</i> , 2013, 4, 104.  | 2.3 | 8         |
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