

# Sara J Margolin

## List of Publications by Year in descending order

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Version: 2024-02-01

57  
papers

4,101  
citations

201674

27  
h-index

155660

55  
g-index

58  
all docs

58  
docs citations

58  
times ranked

8070  
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	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
3	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
4	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
5	<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
6	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
7	E-readers, Computer Screens, or Paper: Does Reading Comprehension Change Across Media Platforms?. <i>Applied Cognitive Psychology</i> , 2013, 27, 512-519.	1.6	113
8	Combined genetic and splicing analysis of BRCA1 c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. <i>Human Molecular Genetics</i> , 2016, 25, 2256-2268.	2.9	106
9	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	12.8	105
10	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.	3.2	94
11	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
12	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
13	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.9	75
14	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	2.4	67
15	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
16	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
17	Fine-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
18	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	2.5	49

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19	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	1.9	45
20	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016, 18, 22.	5.0	43
21	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
22	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	2.9	40
23	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.	3.5	39
24	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
25	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
26	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. <i>Human Molecular Genetics</i> , 2016, 25, 3863-3876.	2.9	33
27	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.	1.8	31
28	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
29	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.	2.5	26
30	<i>Not</i> May Not be Too Difficult: The Effects of Negation on Older Adults' Sentence Comprehension. <i>Educational Gerontology</i> , 2009, 35, 308-322.	1.3	24
31	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
32	CYP2D6 and adjuvant tamoxifen: possible differences of outcome in pre- and post-menopausal patients. <i>Pharmacogenomics</i> , 2013, 14, 613-622.	1.3	21
33	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
34	SNP-SNP interaction analysis of NF- $\kappa$ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , 2015, 6, 37979-37994.	1.8	20
35	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.	3.3	19
36	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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37	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
38	The SNP rs6500843 in 16p13.3 is associated with survival specifically among chemotherapy-treated breast cancer patients. <i>Oncotarget</i> , 2015, 6, 7390-7407.	1.8	15
39	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
40	TP53-based interaction analysis identifies cis-eQTL variants for TP53BP2, FBXO28, and FAM53A that associate with survival and treatment outcome in breast cancer. <i>Oncotarget</i> , 2017, 8, 18381-18398.	1.8	14
41	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
42	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	2.5	12
43	Can Bold Typeface Improve Readers'™ Comprehension and Metacomprehension of Negation?. <i>Reading Psychology</i> , 2013, 34, 85-99.	1.4	11
44	Tumour spectrum in non-BRCA hereditary breast cancer families in Sweden. <i>Hereditary Cancer in Clinical Practice</i> , 2015, 13, 15.	1.5	11
45	Treatment Restarting After Discontinuation of Adjuvant Hormone Therapy in Breast Cancer Patients. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	11
46	Metacomprehension and Negation: Assessing Readers'™ Awareness of the Difficulty of Negated Text. <i>Reading Psychology</i> , 2011, 32, 158-171.	1.4	10
47	Individual Differences in Young and Older Adults' Spelling: Do Good Spellers Age Better than Poor Spellers?. <i>Aging, Neuropsychology, and Cognition</i> , 2007, 14, 529-544.	1.3	9
48	It may not be that difficult the second time around: the effects of rereading on the comprehension and metacomprehension of negated text. <i>Journal of Research in Reading</i> , 2018, 41, 392-402.	2.0	9
49	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	1.8	9
50	How Should I Use My E-Reader? An Exploration of the Circumstances Under Which Electronic Presentation of Text Results in Good Comprehension. <i>Mind, Brain, and Education</i> , 2018, 12, 39-48.	1.9	8
51	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> , 2015, 15, 978.	2.6	6
52	Older Adults'™ Comprehension of Transformational and Deactivation Negation. <i>Educational Gerontology</i> , 2015, 41, 604-612.	1.3	6
53	Cognitively active older adults'™ comprehension and metacomprehension of negated text. <i>Experimental Aging Research</i> , 2018, 44, 329-337.	1.2	1
54	Comprehension and metacomprehension of negated text. <i>Written Language and Literacy</i> , 2020, 23, 92-108.	0.4	1

#	ARTICLE	IF	CITATIONS
55	It's Not Crying, You're Crying: An Evaluation of the Impact of Emotional Text on Negation Comprehension. <i>Reading Psychology</i> , 2021, 42, 131-149.	1.4	1
56	A breast and endometrial cancer syndrome. <i>Maturitas</i> , 2016, 87, 3-4.	2.4	0
57	Parent of Origin and Prognosis in Familial Breast Cancer in Sweden. <i>Anticancer Research</i> , 2017, 37, 1257-1262.	1.1	0