

# Common variants on chromosomes 2q35 and 16q12 confer susceptibility to estrogen receptor- $\alpha$ -positive breast cancer

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Genome-wide association studies of cancer. <i>Future Oncology</i> , 2007, 3, 419-427.	2.4	10
2	Power to Detect Risk Alleles Using Genome-Wide Tag SNP Panels. <i>PLoS Genetics</i> , 2007, 3, e170.	3.5	89
3	Copy Number Variants and Common Disorders: Filling the Gaps and Exploring Complexity in Genome-Wide Association Studies. <i>PLoS Genetics</i> , 2007, 3, e190.	3.5	178
4	Germline genetic variation and breast cancer survival: prognostic and therapeutic implications. <i>Future Oncology</i> , 2007, 3, 491-495.	2.4	4
5	The Genomics Gold Rush. <i>JAMA - Journal of the American Medical Association</i> , 2007, 298, 218.	7.4	71
6	Counterpoint: Genetic Risk Feedback for Common Disease—Time to Test the Waters. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1727-1729.	2.5	7
8	Point: Genetic Risk Feedback for Common Disease—Time to Test the Waters. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1724-1726.	2.5	15
9	Basic science: (MAY 2007). <i>Breast Cancer Online: BCO</i> , 2007, 10, .	0.1	0
10	The promise and limitations of genome-wide association studies to elucidate the causes of breast cancer. <i>Breast Cancer Research</i> , 2007, 9, 114.	5.0	19
11	Genes harbouring susceptibility SNPs are differentially expressed in the breast cancer subtypes. <i>Breast Cancer Research</i> , 2007, 9, 113.	5.0	37
12	Clinical correlates of low-risk variants in FGFR2, TNRC9, MAP3K1, LSP1 and 8q24 in a Dutch cohort of incident breast cancer cases. <i>Breast Cancer Research</i> , 2007, 9, R78.	5.0	64
13	A role for biomarkers in the screening and diagnosis of breast cancer in younger women. <i>Expert Review of Molecular Diagnostics</i> , 2007, 7, 533-544.	3.1	16
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15	Genome-Wide Association Studies: Progress in Identifying Genetic Biomarkers in Common, Complex Diseases. <i>Biomarker Insights</i> , 2007, 2, 117727190700200.	2.5	18
17	Copy number variants and genetic traits: closer to the resolution of phenotypic to genotypic variability. <i>Nature Reviews Genetics</i> , 2007, 8, 639-646.	16.3	391
19	Susceptibility genes in breast cancer: more is less?. <i>Clinical Genetics</i> , 2007, 72, 493-496.	2.0	16
20	Mammary cancer susceptibility: human genes and rodent models. <i>Mammalian Genome</i> , 2007, 18, 817-831.	2.2	32
21	Common Molecular Mechanisms of Mammary Gland Development and Breast Cancer. <i>Cellular and Molecular Life Sciences</i> , 2007, 64, 3185-3200.	5.4	11

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22	Genetic sequence variations and ADPRT haplotype analysis in French Canadian families with high risk of breast cancer. <i>Journal of Human Genetics</i> , 2007, 52, 963-977.	2.3	12
23	The RAD51D E233G variant and breast cancer risk: population-based and clinic-based family studies of Australian women. <i>Breast Cancer Research and Treatment</i> , 2008, 112, 35-39.	2.5	9
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26	Hereditary breast cancer: new genetic developments, new therapeutic avenues. <i>Human Genetics</i> , 2008, 124, 31-42.	3.8	276
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29	Novel breast cancer risk alleles and endometrial cancer risk. <i>International Journal of Cancer</i> , 2008, 123, 2961-2964.	5.1	14
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