

Common variants on chromosomes 2q35 and 16q12 confer susceptibility to estrogen receptor- α 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.	1.1	10
2	Power to Detect Risk Alleles Using Genome-Wide Tag SNP Panels. <i>PLoS Genetics</i> , 2007, 3, e170.	1.5	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.	1.5	178
4	Germline genetic variation and breast cancer survival: prognostic and therapeutic implications. <i>Future Oncology</i> , 2007, 3, 491-495.	1.1	4
5	The Genomics Gold Rush. <i>JAMA - Journal of the American Medical Association</i> , 2007, 298, 218.	3.8	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.	1.1	7
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9	Basic science: (MAY 2007). <i>Breast Cancer Online: BCO</i> , 2007, 10, .	0.1	0
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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.	1.1	9
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#	ARTICLE	IF	CITATIONS
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