

Genome-wide association analysis of more than 120,000 susceptibility loci for breast cancer

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Citation Report

#	ARTICLE	IF	CITATIONS
1	No association between HPV positive breast cancer and expression of human papilloma viral transcripts. <i>Scientific Reports</i> , 2015, 5, 18081.	1.6	21
2	Can the breast screening appointment be used to provide risk assessment and prevention advice?. <i>Breast Cancer Research</i> , 2015, 17, 84.	2.2	30
3	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.	2.2	19
4	Do Variants/Mutations in the CHEK2 Gene Influence the Frequency of Breast and Thyroid Cancer in Certain Families, or the Risk of a Patient Developing Both Thyroid and Breast Cancer?. <i>Clinical Thyroidology</i> , 2015, 27, 89-91.	0.0	0
5	The Role of Constitutional Copy Number Variants in Breast Cancer. <i>Microarrays (Basel, Switzerland)</i> , 2015, 4, 407-423.	1.4	9
6	Managing breast cancer in younger women: challenges and solutions. <i>Breast Cancer: Targets and Therapy</i> , 2016, 8, 1.	1.0	20
7	Tumour characteristics and survival in familial breast cancer prospectively diagnosed by annual mammography. <i>Breast Cancer Research and Treatment</i> , 2015, 152, 87-94.	1.1	2
8	Gene analysis techniques and susceptibility gene discovery in non-BRCA1/BRCA2 familial breast cancer. <i>Surgical Oncology</i> , 2015, 24, 100-109.	0.8	25
9	Invited Commentary: E Pluribus Unum for Epidemiology. <i>American Journal of Epidemiology</i> , 2015, 183, kww236.	1.6	1
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15	The Role of BRCA2 in the Endocytic Trafficking of EGFR and Significance as a Prognostic Biomarker in Cancer. <i>Journal of Cancer</i> , 2016, 7, 2388-2407.	1.2	11
16	Recommendations on Breast Cancer Screening and Prevention in the Context of Implementing Risk Stratification: Impending Changes to Current Policies. <i>Current Oncology</i> , 2016, 23, 615-625.	0.9	37
17	Refining Breast Cancer Risk Stratification: Additional Genes, Additional Information. <i>American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting</i> , 2016, 35, 44-56.	1.8	19
18	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.	0.8	31

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22	Genome-Wide Association Mapping of Acid Soil Resistance in Barley (<i>Hordeum vulgare</i> L.). <i>Frontiers in Plant Science</i> , 2016, 7, 406.	1.7	14
23	Genetic variants in microRNA and microRNA biogenesis pathway genes and breast cancer risk among women of African ancestry. <i>Human Genetics</i> , 2016, 135, 1145-1159.	1.8	32
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148	Prediction of Breast and Prostate Cancer Risks in Male <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers Using Polygenic Risk Scores. <i>Journal of Clinical Oncology</i> , 2017, 35, 2240-2250.	0.8	152
149	Improving radiotherapy in cancer treatment: Promises and challenges. <i>Oncotarget</i> , 2017, 8, 62742-62758.	0.8	186
150	Genetic variation in sensitivity to estrogens and breast cancer risk. <i>Mammalian Genome</i> , 2018, 29, 24-37.	1.0	20
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