

# CITATION REPORT

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**Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes**

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#	Paper	IF	Citations
100	ABC-GWAS: Functional Annotation of Estrogen Receptor-Positive Breast Cancer Genetic Variants. <i>Frontiers in Genetics</i> , <b>2020</b> , 11, 730	4.5	0
99	Update Breast Cancer 2020 Part 3 - Early Breast Cancer. <i>Geburtshilfe Und Frauenheilkunde</i> , <b>2020</b> , 80, 1105-11148		
98	Common variants in signaling transcription-factor-binding sites drive phenotypic variability in red blood cell traits. <i>Nature Genetics</i> , <b>2020</b> , 52, 1333-1345	36.3	8
97	Another step forward towards unraveling the biological mechanisms driving breast cancer predisposition: a role for non-coding RNAs. <i>Non-coding RNA Investigation</i> , <b>2020</b> , 4, 3-3	0.6	
96	The Role of Noncoding Variants in Heritable Disease. <i>Trends in Genetics</i> , <b>2020</b> , 36, 880-891	8.5	25
95	Genetic determinants of breast cancer risk. <i>Current Opinion in Endocrine and Metabolic Research</i> , <b>2020</b> , 15, 1-7	1.7	
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92	Impact of fibroblast growth factor receptor 1 (FGFR1) amplification on the prognosis of breast cancer patients. <i>Breast Cancer Research and Treatment</i> , <b>2020</b> , 184, 311-324	4.4	6
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