

Sarah L Maguire

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

965
citations

840776

11
h-index

996975

15
g-index

16
all docs

16
docs citations

16
times ranked

2837
citing authors

#	ARTICLE	IF	CITATIONS
1	VPRBP Functions Downstream of the Androgen Receptor and OGT to Restrict p53 Activation in Prostate Cancer. <i>Molecular Cancer Research</i> , 2022, 20, 1047-1060.	3.4	2
2	Common Susceptibility Loci for Male Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2021, 113, 453-461.	6.3	12
3	3D Functional Genomics Screens Identify CREBBP as a Targetable Driver in Aggressive Triple-Negative Breast Cancer. <i>Cancer Research</i> , 2021, 81, 847-859.	0.9	7
4	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	6.4	5
5	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203.	6.2	6
6	Capture Hi-C identifies putative target genes at 33 breast cancer risk loci. <i>Nature Communications</i> , 2018, 9, 1028.	12.8	98
7	Evaluation of CDK12 Protein Expression as a Potential Novel Biomarker for DNA Damage Response—Targeted Therapies in Breast Cancer. <i>Molecular Cancer Therapeutics</i> , 2018, 17, 306-315.	4.1	52
8	Three-dimensional modelling identifies novel genetic dependencies associated with breast cancer progression in the isogenic MCF10 model. <i>Journal of Pathology</i> , 2016, 240, 315-328.	4.5	35
9	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	2.9	40
10	SF3B1 mutations constitute a novel therapeutic target in breast cancer. <i>Journal of Pathology</i> , 2015, 235, 571-580.	4.5	167
11	Zinc Finger Transcription Factors Displaced SREBP Proteins as the Major Sterol Regulators during <i>Saccharomyces cerevisiae</i> Evolution. <i>PLoS Genetics</i> , 2014, 10, e1004076.	3.5	63
12	Unbiased analysis of potential targets of breast cancer susceptibility loci by Capture Hi-C. <i>Genome Research</i> , 2014, 24, 1854-1868.	5.5	219
13	Comparative Genome Analysis and Gene Finding in <i>Candida</i> Species Using CGOB. <i>Molecular Biology and Evolution</i> , 2013, 30, 1281-1291.	8.9	115
14	Conserved and Divergent Roles of Bcr1 and CFEM Proteins in <i>Candida parapsilosis</i> and <i>Candida albicans</i> . <i>PLoS ONE</i> , 2011, 6, e28151.	2.5	76
15	Using RNA-seq to determine the transcriptional landscape and the hypoxic response of the pathogenic yeast <i>Candida parapsilosis</i> . <i>BMC Genomics</i> , 2011, 12, 628.	2.8	68