

Sylvie Mazoyer

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

20
papers

940
citations

10
h-index

23
g-index

23
ext. papers

1,132
ext. citations

10
avg, IF

1.57
L-index

#	Paper	IF	Citations
20	Gene- and pathway-level analyses of iCOGS variants highlight novel signaling pathways underlying familial breast cancer susceptibility. <i>International Journal of Cancer</i> , 2021 , 148, 1895-1909	7.5	2
19	Altered regulation of BRCA1 exon 11 splicing is associated with breast cancer risk in carriers of BRCA1 pathogenic variants. <i>Human Mutation</i> , 2021 , 42, 1488-1502	4.7	0
18	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. <i>PLoS ONE</i> , 2020 , 15, e0235655	3.7	1
17	A de novo frameshift pathogenic variant in TBR1 identified in autism without intellectual disability. <i>Human Genomics</i> , 2020 , 14, 32	6.8	2
16	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene 2020 , 15, e0235655		
15	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene 2020 , 15, e0235655		
14	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene 2020 , 15, e0235655		
13	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene 2020 , 15, e0235655		
12	New insights into minor splicing-a transcriptomic analysis of cells derived from TALS patients. <i>Rna</i> , 2019 , 25, 1130-1149	5.8	14
11	Familial breast cancer and DNA repair genes: Insights into known and novel susceptibility genes from the GENESIS study, and implications for multigene panel testing. <i>International Journal of Cancer</i> , 2019 , 144, 1962-1974	7.5	27
10	Novel diagnostic tool for prediction of variant spliceogenicity derived from a set of 395 combined in silico/in vitro studies: an international collaborative effort. <i>Nucleic Acids Research</i> , 2018 , 46, 7913-7923 ^{20.1}		37
9	GEMO, a National Resource to Study Genetic Modifiers of Breast and Ovarian Cancer Risk in and Pathogenic Variant Carriers. <i>Frontiers in Oncology</i> , 2018 , 8, 490	5.3	10
8	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
7	Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , 2017 , 161, 117-134	4.4	15
6	GENESIS: a French national resource to study the missing heritability of breast cancer. <i>BMC Cancer</i> , 2016 , 16, 13	4.8	9
5	Identification of a BRCA2-specific modifier locus at 6p24 related to breast cancer risk. <i>PLoS Genetics</i> , 2013 , 9, e1003173	6	90
4	Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk. <i>PLoS Genetics</i> , 2013 , 9, e1003212	6	209

3	Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. <i>Human Molecular Genetics</i> , 2011 , 20, 4732-47	5.6	21
2	A locus on 19p13 modifies risk of breast cancer in BRCA1 mutation carriers and is associated with hormone receptor-negative breast cancer in the general population. <i>Nature Genetics</i> , 2010 , 42, 885-92	36.3	276
1	Unclassified variants identified in BRCA1 exon 11: Consequences on splicing. <i>Genes Chromosomes and Cancer</i> , 2008 , 47, 418-26	5	33