Sylvie Mazoyer

List of Publications by Citations

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	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. *Nature Genetics*, **2010**, 42, 885-92	36.3	276
19	Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk. *PLoS Genetics*, **2013**, 9, e1003212	6	209
18	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. *Nature Genetics*, **2017**, 49, 1767-1778	36.3	186
17	Identification of a BRCA2-specific modifier locus at 6p24 related to breast cancer risk. *PLoS Genetics*, **2013**, 9, e1003173	6	90
16	Novel diagnostic tool for prediction of variant spliceogenicity derived from a set of 395 combined in silico/in vitro studies: an international collaborative effort. *Nucleic Acids Research*, **2018**, 46, 7913-7923	20.1	37
15	Unclassified variants identified in BRCA1 exon 11: Consequences on splicing. *Genes Chromosomes and Cancer*, **2008**, 47, 418-26	5	33
14	Familial breast cancer and DNA repair genes: Insights into known and novel susceptibility genes from the GENESIS study, and implications for multigene panel testing. *International Journal of Cancer*, **2019**, 144, 1962-1974	7.5	27
13	Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. *Human Molecular Genetics*, **2011**, 20, 4732-47	5.6	21
12	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. *Breast Cancer Research and Treatment*, **2017**, 161, 117-134	4.4	15
11	New insights into minor splicing-a transcriptomic analysis of cells derived from TALS patients. *Rna*, **2019**, 25, 1130-1149	5.8	14
10	GEMO, a National Resource to Study Genetic Modifiers of Breast and Ovarian Cancer Risk in and Pathogenic Variant Carriers. *Frontiers in Oncology*, **2018**, 8, 490	5.3	10
9	GENESIS: a French national resource to study the missing heritability of breast cancer. *BMC Cancer*, **2016**, 16, 13	4.8	9
8	A de novo frameshift pathogenic variant in TBR1 identified in autism without intellectual disability. *Human Genomics*, **2020**, 14, 32	6.8	2
7	Gene- and pathway-level analyses of iCOGS variants highlight novel signaling pathways underlying familial breast cancer susceptibility. *International Journal of Cancer*, **2021**, 148, 1895-1909	7.5	2
6	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. *PLoS ONE*, **2020**, 15, e0235655	3.7	1
5	Altered regulation of BRCA1 exon 11 splicing is associated with breast cancer risk in carriers of BRCA1 pathogenic variants. *Human Mutation*, **2021**, 42, 1488-1502	4.7	0
4	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene **2020**, 15, e0235655		

List of Publications

3 Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene **2020**, 15, e0235655

2 Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene **2020**, 15, e0235655

1 Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene **2020**, 15, e0235655