Sylvie Mazoyer

List of Publications by Year
in descending order

20 papers	1,251 citations	840119 11 h-index	839053 18 g-index
23 all docs	23 docs citations	23 times ranked	3282 citing authors

#	Article	IF	Citations
1	A locus on 19p13 modifies risk of breast cancer in BRCA1 mutation carriers and is associated with hormone receptorâ€“negative breast cancer in the general population. Nature Genetics, 2010, 42, 885-892.	9.4	309
2	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
3	Genome-Wide Association Study in BRCA1 Mutation Carriers Identifies Novel Loci Associated with Breast and Ovarian Cancer Risk. PLoS Genetics, 2013, 9, e1003212.	1.5	244
4	Identification of a BRCA2-Specific Modifier Locus at 6p24 Related to Breast Cancer Risk. PLoS Genetics, 2013, 9, e1003173.	1.5	105
5	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.	6.5	71
6	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.	2.3	50
7	Unclassified variants identified in BRCA1 exon 11: Consequences on splicing. Genes Chromosomes and Cancer, 2008, 47, 418-426.	1.5	36
8	Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. Human Molecular Genetics, 2011, 20, 4732-4747.	1.4	32
9	New insights into minor splicingâ€”a transcriptomic analysis of cells derived from TALS patients. Rna, 2019, 25, 1130-1149.	1.6	27
10	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.	1.1	18
11	GEMO, a National Resource to Study Genetic Modifiers of Breast and Ovarian Cancer Risk in BRCA1 and BRCA2 Pathogenic Variant Carriers. Frontiers in Oncology, 2018, 8, 490.	1.3	14
12	GENESIS: a French national resource to study the missing heritability of breast cancer. BMC Cancer, 2016, 16, 13.	1.1	13
13	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. PLoS ONE, 2020, 15, e0235655.	1.1	8
14	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.	1.1	7
15	A de novo frameshift pathogenic variant in TBR1 identified in autism without intellectual disability. Human Genomics, 2020, 14, 32.	1.4	5
16	Geneâ€•and pathwayâ€level analyses of iCOGS variants highlight novel signaling pathways underlying familial breast cancer susceptibility. International Journal of Cancer, 2021, 148, 1895-1909.	2.3	5
17	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. , 2020, 15, e0235655.		0
18	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. , 2020, 15, e0235655.		0

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19	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. , 2020, 15, e0235655.		0
20	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. , 2020, 15, e0235655.		0