

Michelle W Wong-Brown

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8534743/publications.pdf>

Version: 2024-02-01

19
papers

1,097
citations

758635

12
h-index

794141

19
g-index

21
all docs

21
docs citations

21
times ranked

2399
citing authors

#	ARTICLE	IF	CITATIONS
1	BCL-2 family isoforms in apoptosis and cancer. <i>Cell Death and Disease</i> , 2019, 10, 177.	2.7	394
2	Panel Testing for Familial Breast Cancer: Calibrating the Tension Between Research and Clinical Care. <i>Journal of Clinical Oncology</i> , 2016, 34, 1455-1459.	0.8	154
3	Prevalence of BRCA1 and BRCA2 germline mutations in patients with triple-negative breast cancer. <i>Breast Cancer Research and Treatment</i> , 2015, 150, 71-80.	1.1	103
4	No evidence that protein truncating variants in <i>BRIP1</i> are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016, 53, 298-309.	1.5	94
5	The relative mRNA expression of p53 isoforms in breast cancer is associated with clinical features and outcome. <i>Carcinogenesis</i> , 2014, 35, 586-596.	1.3	67
6	DNA methylation profile of triple negative breast cancer-specific genes comparing lymph node positive patients to lymph node negative patients. <i>Scientific Reports</i> , 2016, 6, 33435.	1.6	56
7	Novel genes associated with lymph node metastasis in triple negative breast cancer. <i>Scientific Reports</i> , 2015, 5, 15832.	1.6	48
8	Prevalence of PALB2 mutations in Australian familial breast cancer cases and controls. <i>Breast Cancer Research</i> , 2015, 17, 111.	2.2	36
9	Reevaluation of the BRCA2 truncating allele c.9976A>&T (p.Lys3326Ter) in a familial breast cancer context. <i>Scientific Reports</i> , 2015, 5, 14800.	1.6	26
10	Mutations in RECQL are not associated with breast cancer risk in an Australian population. <i>Nature Genetics</i> , 2018, 50, 1346-1348.	9.4	19
11	Reevaluation of RINT1 as a breast cancer predisposition gene. <i>Breast Cancer Research and Treatment</i> , 2016, 159, 385-392.	1.1	16
12	The presence of the intron 3 16 bp duplication polymorphism of p53(rs17878362) in breast cancer is associated with a low p53:p53 ratio and better outcome. <i>Carcinogenesis</i> , 2016, 37, 81-86.	1.3	16
13	Epigenetic Mechanisms and Therapeutic Targets in Chemoresistant High-Grade Serous Ovarian Cancer. <i>Cancers</i> , 2021, 13, 5993.	1.7	15
14	Low prevalence of germline <i>PALB2</i> mutations in Australian triple-negative breast cancer. <i>International Journal of Cancer</i> , 2014, 134, 301-305.	2.3	13
15	Investigation of monogenic causes of familial breast cancer: data from the BEACCON case-control study. <i>Npj Breast Cancer</i> , 2021, 7, 76.	2.3	12
16	Methyl-Donor and Cofactor Nutrient Intakes in the First 3 Years and Global DNA Methylation at Age 4: A Prospective Cohort Study. <i>Nutrients</i> , 2018, 10, 273.	1.7	11
17	When is a mutation not a mutation: the case of the c.594-2A>C splice variant in a woman harbouring another BRCA1 mutation in trans. <i>Hereditary Cancer in Clinical Practice</i> , 2016, 14, 6.	0.6	6
18	Global DNA methylation and cognitive and behavioral outcomes at 4 years of age: A cross-sectional study. <i>Brain and Behavior</i> , 2020, 10, e01579.	1.0	6

#	ARTICLE	IF	CITATIONS
19	Sequential azacitidine and carboplatin induces immune activation in platinum-resistant high-grade serous ovarian cancer cell lines and primes for checkpoint inhibitor immunotherapy. BMC Cancer, 2022, 22, 100.	1.1	4