

# Lisa Devereux

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8347053/publications.pdf>

Version: 2024-02-01

23  
papers

609  
citations

759233

12  
h-index

752698

20  
g-index

23  
all docs

23  
docs citations

23  
times ranked

1432  
citing authors

#	ARTICLE	IF	CITATIONS
1	Panel Testing for Familial Breast Cancer: Calibrating the Tension Between Research and Clinical Care. <i>Journal of Clinical Oncology</i> , 2016, 34, 1455-1459.	1.6	154
2	Identification of a new family of human epithelial protein kinases containing two leucine/isoleucine-zipper domains. <i>FEBS Journal</i> , 1993, 213, 701-710.	0.2	107
3	A community-based model of rapid autopsy in end-stage cancer patients. <i>Nature Biotechnology</i> , 2016, 34, 1010-1014.	17.5	66
4	Population-based genetic testing of asymptomatic women for breast and ovarian cancer susceptibility. <i>Genetics in Medicine</i> , 2019, 21, 913-922.	2.4	45
5	Prevalence of PALB2 mutations in Australian familial breast cancer cases and controls. <i>Breast Cancer Research</i> , 2015, 17, 111.	5.0	36
6	The MURAL collection of prostate cancer patient-derived xenografts enables discovery through preclinical models of uro-oncology. <i>Nature Communications</i> , 2021, 12, 5049.	12.8	33
7	Reevaluation of the BRCA2 truncating allele c.9976A>&T (p.Lys3326Ter) in a familial breast cancer context. <i>Scientific Reports</i> , 2015, 5, 14800.	3.3	26
8	Combined Tumor Sequencing and Case-Control Analyses of RAD51C in Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 1332-1338.	6.3	26
9	Evaluating the breast cancer predisposition role of rare variants in genes associated with low-penetrance breast cancer risk SNPs. <i>Breast Cancer Research</i> , 2018, 20, 3.	5.0	19
10	Mutations in RECQL are not associated with breast cancer risk in an Australian population. <i>Nature Genetics</i> , 2018, 50, 1346-1348.	21.4	19
11	Reevaluation of RINT1 as a breast cancer predisposition gene. <i>Breast Cancer Research and Treatment</i> , 2016, 159, 385-392.	2.5	16
12	Molecular comparison of interval and screen-detected breast cancers. <i>Journal of Pathology</i> , 2019, 248, 243-252.	4.5	15
13	Investigation of monogenic causes of familial breast cancer: data from the BEACCON case-control study. <i>Npj Breast Cancer</i> , 2021, 7, 76.	5.2	12
14	A Review of International Biobanks and Networks: Success Factors and Key Benchmarks – A 10-Year Retrospective Review. <i>Biopreservation and Biobanking</i> , 2019, 17, 512-519.	1.0	10
15	Exploring Implementation of Personal Breast Cancer Risk Assessments. <i>Journal of Personalized Medicine</i> , 2021, 11, 992.	2.5	9
16	Evaluation of the association of heterozygous germline variants in NTHL1 with breast cancer predisposition: an international multi-center study of 47,180 subjects. <i>Npj Breast Cancer</i> , 2021, 7, 52.	5.2	7
17	The TP53 mutation rate differs in breast cancers that arise in women with high or low mammographic density. <i>Npj Breast Cancer</i> , 2020, 6, 34.	5.2	4
18	The Clinical and Psychosocial Outcomes for Women Who Received Unexpected Clinically Actionable Germline Information Identified through Research: An Exploratory Sequential Mixed-Methods Comparative Study. <i>Journal of Personalized Medicine</i> , 2022, 12, 1112.	2.5	2

#	ARTICLE	IF	CITATIONS
19	Biobanking in Cancer Research. , 2016, , 27-49.		1
20	Contribution of large genomic rearrangements in <i>PALB2</i> to familial breast cancer: implications for genetic testing. Journal of Medical Genetics, 2023, 60, 112-118.	3.2	1
21	Unselected Women's Experiences of Receiving Genetic Research Results for Hereditary Breast and Ovarian Cancer: A Qualitative Study. Genetic Testing and Molecular Biomarkers, 2021, 25, 741-748.	0.7	1
22	Mixed Lineage Kinases. , 1995, , 369-380.		0
23	Integration of tumour sequencing and case-control data to assess pathogenicity of RAD51C missense variants in familial breast cancer. Npj Breast Cancer, 2022, 8, 10.	5.2	0