

Marike Gabrielson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3914091/publications.pdf>

Version: 2024-02-01

46
papers

4,513
citations

279798

23
h-index

223800

46
g-index

52
all docs

52
docs citations

52
times ranked

7474
citing authors

#	ARTICLE	IF	CITATIONS
1	Prospective evaluation of a breast-cancer risk model integrating classical risk factors and polygenic risk in 15 cohorts from six countries. <i>International Journal of Epidemiology</i> , 2022, 50, 1897-1911.	1.9	43
2	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	4.4	6
3	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744.	7.1	51
4	Circulating proteins reveal prior use of menopausal hormonal therapy and increased risk of breast cancer. <i>Translational Oncology</i> , 2022, 17, 101339.	3.7	1
5	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women. <i>Scientific Reports</i> , 2022, 12, 6199.	3.3	2
6	Topical Endoxifen for Mammographic Density Reduction—A Randomized Controlled Trial. <i>Oncologist</i> , 2022, 27, e597-e600.	3.7	5
7	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51.	8.2	19
8	Distinct Reproductive Risk Profiles for Intrinsic-Like Breast Cancer Subtypes: Pooled Analysis of Population-Based Studies. <i>Journal of the National Cancer Institute</i> , 2022, 114, 1706-1719.	6.3	14
9	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	6.3	45
10	Predictors of mammographic microcalcifications. <i>International Journal of Cancer</i> , 2021, 148, 1132-1143.	5.1	8
11	Breast Cancer Risk Genes — Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	27.0	532
12	Low-Dose Tamoxifen for Mammographic Density Reduction: A Randomized Controlled Trial. <i>Journal of Clinical Oncology</i> , 2021, 39, 1899-1908.	1.6	33
13	Mammographic microcalcifications and risk of breast cancer. <i>British Journal of Cancer</i> , 2021, 125, 759-765.	6.4	32
14	Mammographic Density Change and Risk of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 391-399.	6.3	32
15	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
16	Hormonal determinants of mammographic density and density change. <i>Breast Cancer Research</i> , 2020, 22, 95.	5.0	20
17	Inclusion of Endogenous Plasma Dehydroepiandrosterone Sulfate and Mammographic Density in Risk Prediction Models for Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 574-581.	2.5	6
18	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	1.3	32

#	ARTICLE	IF	CITATIONS
19	PGC1 α and VDAC1 expression in endometrial cancer. <i>Molecular and Clinical Oncology</i> , 2020, 14, 42.	1.0	5
20	Interval breast cancer is associated with other types of tumors. <i>Nature Communications</i> , 2019, 10, 4648.	12.8	25
21	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	5.2	28
22	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	3.3	5
23	Determinants of Mammographic Density Change. <i>JNCI Cancer Spectrum</i> , 2019, 3, pkz004.	2.9	27
24	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	12.8	90
25	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	6.4	52
26	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	6.2	711
27	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 795-806.	1.9	81
28	The <i>BRCA2</i> c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018, 39, 729-741.	2.5	19
29	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. <i>International Journal of Epidemiology</i> , 2018, 47, 526-536.	1.9	88
30	Affinity proteomic profiling of plasma for proteins associated to area-based mammographic breast density. <i>Breast Cancer Research</i> , 2018, 20, 14.	5.0	8
31	Association of reproductive history with breast tissue characteristics and receptor status in the normal breast. <i>Breast Cancer Research and Treatment</i> , 2018, 170, 487-497.	2.5	15
32	Inclusion of Plasma Prolactin Levels in Current Risk Prediction Models of Premenopausal and Postmenopausal Breast Cancer. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky055.	2.9	16
33	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	21.4	184
34	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017, 49, 834-841.	21.4	426
35	Cohort Profile: The Karolinska Mammography Project for Risk Prediction of Breast Cancer (KARMA). <i>International Journal of Epidemiology</i> , 2017, 46, 1740-1741g.	1.9	88
36	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099

#	ARTICLE	IF	CITATIONS
37	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
38	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	1.8	9
39	Amount of stroma is associated with mammographic density and stromal expression of oestrogen receptor in normal breast tissues. <i>Breast Cancer Research and Treatment</i> , 2016, 158, 253-261.	2.5	17
40	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	12.8	93
41	Altered PPAR γ Coactivator-1 Alpha Expression in Abdominal Aortic Aneurysm: Possible Effects on Mitochondrial Biogenesis. <i>Journal of Vascular Research</i> , 2016, 53, 17-26.	1.4	15
42	Mitochondrial regulation of cell cycle progression through SLC25A43. <i>Biochemical and Biophysical Research Communications</i> , 2016, 469, 1090-1096.	2.1	11
43	Expression of Mitochondrial Regulators PGC1 α and TFAM as Putative Markers of Subtype and Chemoresistance in Epithelial Ovarian Carcinoma. <i>PLoS ONE</i> , 2014, 9, e107109.	2.5	35
44	The mitochondrial transport protein SLC25A43 affects drug efficacy and drug-induced cell cycle arrest in breast cancer cell lines. <i>Oncology Reports</i> , 2013, 29, 1268-1274.	2.6	7
45	The mitochondrial transporter SLC25A43 is frequently deleted and may influence cell proliferation in HER2-positive breast tumors. <i>BMC Cancer</i> , 2012, 12, 350.	2.6	16
46	Inhibition of Hedgehog Signaling Decreases Proliferation and Clonogenicity of Human Mesenchymal Stem Cells. <i>PLoS ONE</i> , 2011, 6, e16798.	2.5	47