

# Julika Borde

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

4,295  
citations

394286

19  
h-index

526166

27  
g-index

28  
all docs

28  
docs citations

28  
times ranked

6844  
citing authors

#	ARTICLE	IF	CITATIONS
1	Breast and Prostate Cancer Risks for Male <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variant Carriers Using Polygenic Risk Scores. <i>Journal of the National Cancer Institute</i> , 2022, 114, 109-122.	3.0	19
2	Clonal Hematopoiesis-Associated Gene Mutations in a Clinical Cohort of 448 Patients With Ovarian Cancer. <i>Journal of the National Cancer Institute</i> , 2022, 114, 565-570.	3.0	17
3	A case-only study to identify genetic modifiers of breast cancer risk for <i>BRCA1/BRCA2</i> mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	5.8	19
4	Breast Cancer Risk Genes Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	13.9	532
5	Performance of In Silico Prediction Tools for the Detection of Germline Copy Number Variations in Cancer Predisposition Genes in 4208 Female Index Patients with Familial Breast and Ovarian Cancer. <i>Cancers</i> , 2021, 13, 118.	1.7	4
6	Performance of Breast Cancer Polygenic Risk Scores in 760 Female <i>CHEK2</i> Germline Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2021, 113, 893-899.	3.0	21
7	Association of Genomic Domains in <i>BRCA1</i> and <i>BRCA2</i> with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , 2020, 80, 624-638.	0.4	39
8	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	1.1	82
9	Characterization of the Cancer Spectrum in Men With Germline <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>JAMA Oncology</i> , 2020, 6, 1218.	3.4	48
10	Ovarian and Breast Cancer Risks Associated With Pathogenic Variants in <i>RAD51C</i> and <i>RAD51D</i> . <i>Journal of the National Cancer Institute</i> , 2020, 112, 1242-1250.	3.0	106
11	Germline loss-of-function variants in the <i>BARD1</i> gene are associated with early-onset familial breast cancer but not ovarian cancer. <i>Breast Cancer Research</i> , 2019, 21, 55.	2.2	44
12	Deleterious somatic variants in 473 consecutive individuals with ovarian cancer: results of the observational AGO-TR1 study (NCT02222883). <i>Journal of Medical Genetics</i> , 2019, 56, 574-580.	1.5	34
13	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	2.6	711
14	The <i>GPRC5A</i> frameshift variant c.183del is not associated with increased breast cancer risk in <i>BRCA1</i> mutation carriers. <i>International Journal of Cancer</i> , 2019, 144, 1761-1763.	2.3	2
15	Gene panel testing of 5589 <i>BRCA1/2</i> -negative index patients with breast cancer in a routine diagnostic setting: results of the German Consortium for Hereditary Breast and Ovarian Cancer. <i>Cancer Medicine</i> , 2018, 7, 1349-1358.	1.3	126
16	Diagnosis of Li-Fraumeni Syndrome: Differentiating <i>TP53</i> germline mutations from clonal hematopoiesis. <i>Human Mutation</i> , 2018, 39, 2040-2046.	1.1	20
17	<i>BRIP1</i> loss-of-function mutations confer high risk for familial ovarian cancer, but not familial breast cancer. <i>Breast Cancer Research</i> , 2018, 20, 7.	2.2	78
18	Germline Mutations in Triple-Negative Breast Cancer. <i>Breast Care</i> , 2017, 12, 15-19.	0.8	59

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19	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
20	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
21	Prevalence of deleterious germline variants in risk genes including BRCA1/2 in consecutive ovarian cancer patients (AGO-TR-1). <i>PLoS ONE</i> , 2017, 12, e0186043.	1.1	105
22	Prediction of Breast and Prostate Cancer Risks in Male <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers Using Polygenic Risk Scores. <i>Journal of Clinical Oncology</i> , 2017, 35, 2240-2250.	0.8	152
23	Prevalence of <i>BRCA1/2</i> germline mutations in 21â€¦401 families with breast and ovarian cancer. <i>Journal of Medical Genetics</i> , 2016, 53, 465-471.	1.5	179
24	Male breast cancer in <i>BRCA1</i> and <i>BRCA2</i> mutation carriers: pathology data from the Consortium of Investigators of Modifiers of <i>BRCA1/2</i> . <i>Breast Cancer Research</i> , 2016, 18, 15.	2.2	88
25	Association of Type and Location of <i>BRCA1</i> and <i>BRCA2</i> Mutations With Risk of Breast and Ovarian Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1347.	3.8	390
26	<i>RAD51C</i> deletion screening identifies a recurrent gross deletion in breast cancer and ovarian cancer families. <i>Breast Cancer Research</i> , 2013, 15, R120.	2.2	28