

# Marinus J Blok

## List of Publications by Year in descending order

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Version: 2024-02-01

25  
papers

2,030  
citations

394421

19  
h-index

580821

25  
g-index

25  
all docs

25  
docs citations

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times ranked

4755  
citing authors

#	ARTICLE	IF	CITATIONS
1	Prevalence of Germline Pathogenic Variants in Cancer Predisposing Genes in Czech and Belgian Pancreatic Cancer Patients. <i>Cancers</i> , 2021, 13, 4430.	3.7	8
2	Pathogenic neurofibromatosis type 1 (NF1) RNA splicing resolved by targeted RNAseq. <i>Npj Genomic Medicine</i> , 2021, 6, 95.	3.8	9
3	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.	0.9	39
4	Generation and initial characterization of novel tumour organoid models to study human pancreatic cancer-induced cachexia. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2020, 11, 1509-1524.	7.3	29
5	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.	2.4	82
6	Alcohol Consumption, Cigarette Smoking, and Risk of Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Results from The <i>BRCA1</i> and <i>BRCA2</i> Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 368-378.	2.5	24
7	Dutch genome diagnostic laboratories accelerated and improved variant interpretation and increased accuracy by sharing data. <i>Human Mutation</i> , 2019, 40, 2230-2238.	2.5	32
8	Survival after bilateral risk-reducing mastectomy in healthy <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2019, 177, 723-733.	2.5	111
9	Targeted RNAseq successfully identifies normal and pathogenic splicing events in breast/ovarian cancer susceptibility and Lynch syndrome genes. <i>International Journal of Cancer</i> , 2019, 145, 401-414.	5.1	27
10	Mutational spectrum in a worldwide study of 29,700 families with <i>BRCA1</i> or <i>BRCA2</i> mutations. <i>Human Mutation</i> , 2018, 39, 593-620.	2.5	224
11	The <i>BRCA1</i> c. 5096G>A p.Arg1699Gln (R1699Q) intermediate risk variant: breast and ovarian cancer risk estimation and recommendations for clinical management from the ENIGMA consortium. <i>Journal of Medical Genetics</i> , 2018, 55, 15-20.	3.2	50
12	<i>BRCA1</i> and <i>BRCA2</i> 5' noncoding region variants identified in breast cancer patients alter promoter activity and protein binding. <i>Human Mutation</i> , 2018, 39, 2025-2039.	2.5	15
13	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	21.4	356
14	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
15	Association of breast cancer risk in <i>BRCA1</i> and <i>BRCA2</i> mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 117-134.	2.5	18
16	Novel <i>BRCA1</i> and <i>BRCA2</i> Tumor Test as Basis for Treatment Decisions and Referral for Genetic Counselling of Patients with Ovarian Carcinomas. <i>Human Mutation</i> , 2017, 38, 226-235.	2.5	55
17	Next-Generation Sequencing in Oncology: Genetic Diagnosis, Risk Prediction and Cancer Classification. <i>International Journal of Molecular Sciences</i> , 2017, 18, 308.	4.1	353
18	Naturally occurring <i>BRCA2</i> alternative mRNA splicing events in clinically relevant samples. <i>Journal of Medical Genetics</i> , 2016, 53, 548-558.	3.2	69

#	ARTICLE	IF	CITATIONS
19	Growth pattern in Kabuki syndrome with a <i>KMT2D</i> mutation. American Journal of Medical Genetics, Part A, 2016, 170, 3172-3179.	1.2	23
20	Assessing Associations between the AURKA-HMMR-TPX2-TUBG1 Functional Module and Breast Cancer Risk in BRCA1/2 Mutation Carriers. PLoS ONE, 2015, 10, e0120020.	2.5	34
21	Germ-line variants identified by next generation sequencing in a panel of estrogen and cancer associated genes correlate with poor clinical outcome in Lynch syndrome patients. Oncotarget, 2015, 6, 41108-41122.	1.8	5
22	BRCA1 Circos: a visualisation resource for functional analysis of missense variants. Journal of Medical Genetics, 2015, 52, 224-230.	3.2	32
23	Comprehensive annotation of splice junctions supports pervasive alternative splicing at the BRCA1 locus: a report from the ENIGMA consortium. Human Molecular Genetics, 2014, 23, 3666-3680.	2.9	96
24	PGD for hereditary breast and ovarian cancer: the route to universal tests for BRCA1 and BRCA2 mutation carriers. European Journal of Human Genetics, 2013, 21, 1361-1368.	2.8	21
25	Characterisation of unclassified variants in the BRCA1/2 genes with a putative effect on splicing. Breast Cancer Research and Treatment, 2011, 129, 971-982.	2.5	29