

# Hanne Meijers-Heijboer

## List of Publications by Year in descending order

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Version: 2024-02-01

30  
papers

3,973  
citations

430874

18  
h-index

526287

27  
g-index

30  
all docs

30  
docs citations

30  
times ranked

8374  
citing authors

#	ARTICLE	IF	CITATIONS
1	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	21.4	960
3	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	21.4	513
4	Swarm Intelligence-Enhanced Detection of Non-Small-Cell Lung Cancer Using Tumor-Educated Platelets. <i>Cancer Cell</i> , 2017, 32, 238-252.e9.	16.8	235
5	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
6	A novel Fanconi anaemia subtype associated with a dominant-negative mutation in RAD51. <i>Nature Communications</i> , 2015, 6, 8829.	12.8	130
7	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	8.4	118
8	Characterization of pathogenic SORL1 genetic variants for association with Alzheimer's disease: a clinical interpretation strategy. <i>European Journal of Human Genetics</i> , 2017, 25, 973-981.	2.8	102
9	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
10	Somatic genomic alterations in retinoblastoma beyond RB1 are rare and limited to copy number changes. <i>Scientific Reports</i> , 2016, 6, 25264.	3.3	75
11	RB1 mutation spectrum in a comprehensive nationwide cohort of retinoblastoma patients. <i>Journal of Medical Genetics</i> , 2014, 51, 366-374.	3.2	69
12	A Meta-Analysis of Retinoblastoma Copy Numbers Refines the List of Possible Driver Genes Involved in Tumor Progression. <i>PLoS ONE</i> , 2016, 11, e0153323.	2.5	55
13	Loss of photoreceptor and gain of genomic alterations in retinoblastoma reveal tumor progression. <i>EBioMedicine</i> , 2015, 2, 660-670.	6.1	54
14	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	2.5	49
15	Genomic profiling of CHEK2*1100delC-mutated breast carcinomas. <i>BMC Cancer</i> , 2015, 15, 877.	2.6	44
16	Identification of a Dutch founder mutation in MUSK causing fetal akinesia deformation sequence. <i>European Journal of Human Genetics</i> , 2015, 23, 1151-1157.	2.8	42
17	Addition of a 161-SNP polygenic risk score to family history-based risk prediction: impact on clinical management in non-BRCA1/2 breast cancer families. <i>Journal of Medical Genetics</i> , 2019, 56, 581-589.	3.2	35
18	Targeted carrier screening for four recessive disorders: High detection rate within a founder population. <i>European Journal of Medical Genetics</i> , 2015, 58, 123-128.	1.3	31

#	ARTICLE	IF	CITATIONS
19	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016, 27, 679-693.	1.8	21
20	Proper genomic profiling of ( <i>BRCA1</i> mutated) basal-like breast carcinomas requires prior removal of tumor infiltrating lymphocytes. <i>Molecular Oncology</i> , 2015, 9, 877-888.	4.6	16
21	Uptake of prenatal diagnostic testing for retinoblastoma compared to other hereditary cancer syndromes in the Netherlands. <i>Familial Cancer</i> , 2017, 16, 271-277.	1.9	16
22	First steps in exploring prospective exome sequencing of consanguineous couples. <i>European Journal of Medical Genetics</i> , 2014, 57, 613-616.	1.3	11
23	The Effect of Predictive Testing in Adult-Onset Neurodegenerative Diseases on Social and Personal Life. <i>Journal of Genetic Counseling</i> , 2018, 27, 947-954.	1.6	10
24	A novel <i>CCM2</i> variant in a family with non-progressive cognitive complaints and cerebral microbleeds. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 220-226.	1.7	6
25	Coregulation of <i>FANCA</i> and <i>BRCA1</i> in human cells. <i>SpringerPlus</i> , 2014, 3, 381.	1.2	5
26	Genomic landscape of retinoblastoma in <i>Rb<sup>+/+</sup>p130<sup>+/+</sup></i> mice resembles human retinoblastoma. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 231-242.	2.8	5
27	Direct-to-consumer carrier screening for cystic fibrosis via a hospital website: a 6-year evaluation. <i>Journal of Community Genetics</i> , 2019, 10, 249-257.	1.2	5
28	S4-01-01: Pathology of cognitively healthy centenarians. , 2015, 11, P257-P257.		2
29	P2-134: THE ADDED VALUE OF EXTREME PHENOTYPES IN ALZHEIMER'S DISEASE CASE-CONTROL STUDIES. <i>Alzheimer's and Dementia</i> , 2018, 14, P719.	0.8	0
30	P4-038: IS <i>SORL1</i> AN AUTOSOMAL DOMINANT ALZHEIMER GENE?. <i>Alzheimer's and Dementia</i> , 2018, 14, P1447.	0.8	0