

Christi J Van Asperen

List of Publications by Year in descending order

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Version: 2024-02-01

155
papers

11,898
citations

36203

51
h-index

31759

101
g-index

159
all docs

159
docs citations

159
times ranked

15564
citing authors

#	ARTICLE	IF	CITATIONS
1	Universal Immunohistochemistry for Lynch Syndrome: A Systematic Review and Meta-analysis of 58,580 Colorectal Carcinomas. <i>Clinical Gastroenterology and Hepatology</i> , 2022, 20, e496-e507.	2.4	14
2	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
3	Risks of breast and ovarian cancer for women harboring pathogenic missense variants in <i>BRCA1</i> and <i>BRCA2</i> compared with those harboring protein truncating variants. <i>Genetics in Medicine</i> , 2022, 24, 119-129.	1.1	10
4	Effects of chemotherapy on contralateral breast cancer risk in <i>BRCA1</i> and <i>BRCA2</i> mutation carriers: A nationwide cohort study. <i>Breast</i> , 2022, 61, 98-107.	0.9	6
5	Recommendations for reporting results of diagnostic genomic testing. <i>European Journal of Human Genetics</i> , 2022, 30, 1011-1016.	1.4	15
6	Genetic clinicians' confidence in <i>BOADICEA</i> comprehensive breast cancer risk estimates and counselees' psychosocial outcomes: A prospective study. <i>Clinical Genetics</i> , 2022, 102, 30-39.	1.0	3
7	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51.	3.6	19
8	Association between a 46-SNP Polygenic Risk Score and melanoma risk in Dutch patients with familial melanoma. <i>Journal of Medical Genetics</i> , 2021, 58, 760-766.	1.5	8
9	Reproductive decision-making in the context of hereditary cancer: the effects of an online decision aid on informed decision-making. <i>Journal of Community Genetics</i> , 2021, 12, 101-110.	0.5	6
10	Long-Term Morbidity and Health After Early Menopause Due to Oophorectomy in Women at Increased Risk of Ovarian Cancer: Protocol for a Nationwide Cross-Sectional Study With Prospective Follow-Up (HARMONY Study). <i>JMIR Research Protocols</i> , 2021, 10, e24414.	0.5	9
11	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
12	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
13	Endometrial Cancer Risk in Women With Germline <i>BRCA1</i> or <i>BRCA2</i> Mutations: Multicenter Cohort Study. <i>Journal of the National Cancer Institute</i> , 2021, 113, 1203-1211.	3.0	44
14	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous <i>BRCA1</i> or <i>BRCA2</i> pathogenic variant. <i>Genetics in Medicine</i> , 2021, 23, 1726-1737.	1.1	16
15	Evaluation of multiple transcriptomic gene risk signatures in male breast cancer. <i>Npj Breast Cancer</i> , 2021, 7, 98.	2.3	4
16	Response to Nahshon and Lavie. <i>Journal of the National Cancer Institute</i> , 2021, , .	3.0	0
17	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
18	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120

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19	Clustering of known low and moderate risk alleles rather than a novel recessive high-risk gene in non-BRCA1/2 sib trios affected with breast cancer. <i>International Journal of Cancer</i> , 2020, 147, 2708-2716.	2.3	2
20	Alternative mRNA splicing can attenuate the pathogenicity of presumed loss-of-function variants in BRCA2. <i>Genetics in Medicine</i> , 2020, 22, 1355-1365.	1.1	23
21	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	1.6	2
22	Characterization of the Cancer Spectrum in Men With Germline BRCA1 and BRCA2 Pathogenic Variants. <i>JAMA Oncology</i> , 2020, 6, 1218.	3.4	48
23	–Do Preferred Risk Formats Lead to Better Understanding? A Multicenter Controlled Trial on Communicating Familial Breast Cancer Risks Using Different Risk Formats–. <i>Patient Preference and Adherence</i> , 2020, Volume 14, 333-342.	0.8	4
24	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
25	Abstract P4-10-03: The genomic landscape of male breast cancers using the oncomine comprehensive assay for actionable mutations. , 2020, , .		0
26	The functional impact of variants of uncertain significance in BRCA2. <i>Genetics in Medicine</i> , 2019, 21, 293-302.	1.1	58
27	Survival after bilateral risk-reducing mastectomy in healthy BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2019, 177, 723-733.	1.1	111
28	Breast Cancer Susceptibility–Towards Individualised Risk Prediction. <i>Current Genetic Medicine Reports</i> , 2019, 7, 124-135.	1.9	4
29	Interim Results from the IMPACT Study: Evidence for Prostate-specific Antigen Screening in BRCA2 Mutation Carriers. <i>European Urology</i> , 2019, 76, 831-842.	0.9	148
30	Large scale multifactorial likelihood quantitative analysis of BRCA1 and BRCA2 variants: An ENIGMA resource to support clinical variant classification. <i>Human Mutation</i> , 2019, 40, 1557-1578.	1.1	102
31	An alternative approach to establishing unbiased colorectal cancer risk estimation in Lynch syndrome. <i>Genetics in Medicine</i> , 2019, 21, 2706-2712.	1.1	11
32	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	2.9	19
33	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.	1.5	35
34	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
35	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	2.9	52
36	Germline BRCA-Associated Endometrial Carcinoma Is a Distinct Clinicopathologic Entity. <i>Clinical Cancer Research</i> , 2019, 25, 7517-7526.	3.2	34

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37	Multigene panel sequencing of established and candidate melanoma susceptibility genes in a large cohort of Dutch non-CDKN2A/CDK4 melanoma families. <i>International Journal of Cancer</i> , 2019, 144, 2453-2464.	2.3	33
38	Online decision support for persons having a genetic predisposition to cancer and their partners during reproductive decision-making. <i>Journal of Genetic Counseling</i> , 2019, 28, 533-542.	0.9	14
39	Height and Body Mass Index as Modifiers of Breast Cancer Risk in BRCA1/2 Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019, 111, 350-364.	3.0	30
40	A functional assay-based procedure to classify mismatch repair gene variants in Lynch syndrome. <i>Genetics in Medicine</i> , 2019, 21, 1486-1496.	1.1	36
41	The development of an online decision aid to support persons having a genetic predisposition to cancer and their partners during reproductive decision-making: a usability and pilot study. <i>Familial Cancer</i> , 2019, 18, 137-146.	0.9	17
42	The BRCA2 c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018, 39, 729-741.	1.1	19
43	CM-Score: a validated scoring system to predict CDKN2A germline mutations in melanoma families from Northern Europe. <i>Journal of Medical Genetics</i> , 2018, 55, 661-668.	1.5	13
44	Mutational spectrum in a worldwide study of 29,700 families with BRCA1 or BRCA2 mutations. <i>Human Mutation</i> , 2018, 39, 593-620.	1.1	224
45	Prostate-specific antigen velocity in a prospective prostate cancer screening study of men with genetic predisposition. <i>British Journal of Cancer</i> , 2018, 118, 266-276.	2.9	12
46	E-cadherin breast tumor expression, risk factors and survival: Pooled analysis of 5,933 cases from 12 studies in the Breast Cancer Association Consortium. <i>Scientific Reports</i> , 2018, 8, 6574.	1.6	51
47	Oral Contraceptive Use and Breast Cancer Risk: Retrospective and Prospective Analyses From a BRCA1 and BRCA2 Mutation Carrier Cohort Study. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky023.	1.4	33
48	Ovarian stimulation for IVF and risk of primary breast cancer in BRCA1/2 mutation carriers. <i>British Journal of Cancer</i> , 2018, 119, 357-363.	2.9	22
49	Validation and Implementation of BRCA1/2 Variant Screening in Ovarian Tumor Tissue. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 600-611.	1.2	18
50	Male breast cancer precursor lesions: analysis of the EORTC 10085/TBCRC/BIG/NABCG International Male Breast Cancer Program. <i>Modern Pathology</i> , 2017, 30, 509-518.	2.9	32
51	BRCA2 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.4	75
52	Pathological characterisation of male breast cancer: Results of the EORTC 10085/TBCRC/BIG/NABCG International Male Breast Cancer Program. <i>European Journal of Cancer</i> , 2017, 82, 219-227.	1.3	71
53	Guidance Statement On BRCA1/2 Tumor Testing in Ovarian Cancer Patients. <i>Seminars in Oncology</i> , 2017, 44, 187-197.	0.8	76
54	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099

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55	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
56	Validation of a scale for assessing attitudes towards outcomes of genetic cancer testing among primary care providers and breast specialists. <i>PLoS ONE</i> , 2017, 12, e0178447.	1.1	9
57	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
58	Functional Analysis of Missense Variants in the Putative Breast Cancer Susceptibility Gene <i>XRCC2</i> . <i>Human Mutation</i> , 2016, 37, 914-925.	1.1	12
59	Germline <i>BRCA1/2</i> mutation testing is indicated in every patient with epithelial ovarian cancer: A systematic review. <i>European Journal of Cancer</i> , 2016, 61, 137-145.	1.3	64
60	Comprehensive Mutation Analysis of <i>PMS2</i> in a Large Cohort of Proband Suspected of Lynch Syndrome or Constitutional Mismatch Repair Deficiency Syndrome. <i>Human Mutation</i> , 2016, 37, 1162-1179.	1.1	50
61	Current perspectives on recommendations for <i>BRCA</i> genetic testing in ovarian cancer patients. <i>European Journal of Cancer</i> , 2016, 69, 127-134.	1.3	49
62	Do <i>BRCA1/2</i> mutation carriers have an earlier onset of natural menopause?. <i>Menopause</i> , 2016, 23, 903-910.	0.8	22
63	Bias Explains Most of the Parent-of-Origin Effect on Breast Cancer Risk in <i>BRCA1/2</i> Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1251-1258.	1.1	9
64	Classification and Clinical Management of Variants of Uncertain Significance in High Penetrance Cancer Predisposition Genes. <i>Human Mutation</i> , 2016, 37, 331-336.	1.1	31
65	Combined genetic and splicing analysis of <i>BRCA1</i> c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. <i>Human Molecular Genetics</i> , 2016, 25, 2256-2268.	1.4	106
66	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	428
67	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	1.4	40
68	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating <i>MAP3K1</i> . <i>American Journal of Human Genetics</i> , 2015, 96, 5-20.	2.6	76
69	Inherited variants in the inner centromere protein (<i>INCENP</i>) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. <i>Carcinogenesis</i> , 2015, 36, 256-271.	1.3	14
70	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.	9.4	513
71	The role of the prostate cancer gene 3 urine test in addition to serum prostate-specific antigen level in prostate cancer screening among breast cancer, early-onset gene mutation carriers. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2015, 33, 202.e19-202.e28.	0.8	8
72	<i>FANCM</i> c.5791C>T nonsense mutation (rs144567652) induces exon skipping, affects DNA repair activity and is a familial breast cancer risk factor. <i>Human Molecular Genetics</i> , 2015, 24, 5345-5355.	1.4	91

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73	General Practitioners and Breast Surgeons in France, Germany, Netherlands and the UK show variable breast cancer risk communication profiles. <i>BMC Cancer</i> , 2015, 15, 243.	1.1	6
74	Next-Generation Diagnostics: Gene Panel, Exome, or Whole Genome?. <i>Human Mutation</i> , 2015, 36, 648-655.	1.1	124
75	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv219.	3.0	99
76	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 285-298.	1.4	38
77	Abstract S6-05: Characterization of male breast cancer: First results of the EORTC10085/TBCRC/BIG/NABCG International Male BC Program. , 2015, , .		20
78	Breast surgeons' attitudes towards bilateral risk-reducing mastectomy: A National Survey of American Surgeons.. <i>Journal of Clinical Oncology</i> , 2015, 33, 25-25.	0.8	3
79	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	1.1	49
80	DNA Glycosylases Involved in Base Excision Repair May Be Associated with Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>PLoS Genetics</i> , 2014, 10, e1004256.	1.5	47
81	Meaning-centered group psychotherapy in cancer survivors: a feasibility study. <i>Psycho-Oncology</i> , 2014, 23, 827-831.	1.0	19
82	Refined histopathological predictors of BRCA1 and BRCA2 mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. <i>Breast Cancer Research</i> , 2014, 16, 3419.	2.2	97
83	Breast and ovarian cancer risks in a large series of clinically ascertained families with a high proportion of BRCA1 and BRCA2 Dutch founder mutations. <i>Journal of Medical Genetics</i> , 2014, 51, 98-107.	1.5	74
84	Variation in Mutation Spectrum Partly Explains Regional Differences in the Breast Cancer Risk of Female BRCA Mutation Carriers in the Netherlands. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2482-2491.	1.1	11
85	Cancer risk communication, predictive testing and management in France, Germany, the Netherlands and the UK: general practitioners' and breast surgeons' current practice and preferred practice responsibilities. <i>Journal of Community Genetics</i> , 2014, 5, 69-79.	0.5	16
86	Targeted Prostate Cancer Screening in BRCA1 and BRCA2 Mutation Carriers: Results from the Initial Screening Round of the IMPACT Study. <i>European Urology</i> , 2014, 66, 489-499.	0.9	195
87	Relevance and efficacy of breast cancer screening in BRCA1 and BRCA2 mutation carriers above 60 years: A national cohort study. <i>International Journal of Cancer</i> , 2014, 135, 2940-2949.	2.3	13
88	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
89	Effectiveness and cost-effectiveness of meaning-centered group psychotherapy in cancer survivors: protocol of a randomized controlled trial. <i>BMC Psychiatry</i> , 2014, 14, 22.	1.1	44
90	CHEK2*1100delC homozygosity in the Netherlands prevalence and risk of breast and lung cancer. <i>European Journal of Human Genetics</i> , 2014, 22, 46-51.	1.4	29

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91	The counselees' self-reported request for psychological help in genetic counseling for hereditary breast/ovarian cancer: not only psychopathology matters. <i>Psycho-Oncology</i> , 2013, 22, 902-910.	1.0	35
92	Genetic counseling does not fulfill the counselees' need for certainty in hereditary breast/ovarian cancer families: an explorative assessment. <i>Psycho-Oncology</i> , 2013, 22, 1167-1176.	1.0	19
93	The effectiveness of a graphical presentation in addition to a frequency format in the context of familial breast cancer risk communication: a multicenter controlled trial. <i>BMC Medical Informatics and Decision Making</i> , 2013, 13, 55.	1.5	10
94	Value-based healthcare in Lynch syndrome. <i>Familial Cancer</i> , 2013, 12, 347-354.	0.9	6
95	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
96	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
97	Variants of Uncertain Significance in <i>BRCA1</i> and <i>BRCA2</i> assessment of in silico analysis and a proposal for communication in genetic counselling. <i>Journal of Medical Genetics</i> , 2013, 50, 74-79.	1.5	21
98	Whole Exome Sequencing Suggests Much of Non- <i>BRCA1/BRCA2</i> Familial Breast Cancer Is Due to Moderate and Low Penetrance Susceptibility Alleles. <i>PLoS ONE</i> , 2013, 8, e55681.	1.1	95
99	Efficacy of risk-reducing mastectomy (RRM) on overall survival (OS) in <i>BRCA1/2</i> -associated breast cancer (BC) patients.. <i>Journal of Clinical Oncology</i> , 2013, 31, 1502-1502.	0.8	3
100	Exome Sequencing of Germline DNA from Non- <i>BRCA1/2</i> Familial Breast Cancer Cases Selected on the Basis of aCGH Tumor Profiling. <i>PLoS ONE</i> , 2013, 8, e55734.	1.1	29
101	A Nonsynonymous Polymorphism in <i>IRS1</i> Modifies Risk of Developing Breast and Ovarian Cancers in <i>BRCA1</i> and Ovarian Cancer in <i>BRCA2</i> Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1362-1370.	1.1	23
102	<i>BRCA1</i> R1699Q variant displaying ambiguous functional abrogation confers intermediate breast and ovarian cancer risk. <i>Journal of Medical Genetics</i> , 2012, 49, 525-532.	1.5	97
103	Exposure to diagnostic radiation and risk of breast cancer among carriers of <i>BRCA1/2</i> mutations: retrospective cohort study (GENE-RAD-RISK). <i>BMJ</i> , The, 2012, 345, e5660-e5660.	3.0	186
104	Rare variants in <i>XRCC2</i> as breast cancer susceptibility alleles: Table 1. <i>Journal of Medical Genetics</i> , 2012, 49, 618-620.	1.5	49
105	Breast Cancer Risk and 6q22.33: Combined Results from Breast Cancer Association Consortium and Consortium of Investigators on Modifiers of <i>BRCA1/2</i> . <i>PLoS ONE</i> , 2012, 7, e35706.	1.1	11
106	Opening the psychological black box in genetic counseling. The psychological impact of DNA testing is predicted by the counselees' perception, the medical impact by the pathogenic or uninformative <i>BRCA1/2</i> result. <i>Psycho-Oncology</i> , 2012, 21, 29-42.	1.0	60
107	Urologists' and GPs' knowledge of hereditary prostate cancer is suboptimal for prostate cancer counseling: a nation-wide survey in The Netherlands. <i>Familial Cancer</i> , 2012, 11, 195-200.	0.9	6
108	<i>MUTYH</i> gene variants and breast cancer in a Dutch case-control study. <i>Breast Cancer Research and Treatment</i> , 2012, 134, 219-227.	1.1	38

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109	Exploring the short-term impact of DNA-testing in breast cancer patients: The counselees' perception matters, but the actual BRCA1/2 result does not. <i>Patient Education and Counseling</i> , 2012, 86, 239-251.	1.0	31
110	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 3289-3303.	1.4	152
111	A counselee-oriented perspective on risk communication in genetic counseling: Explaining the inaccuracy of the counselees' risk perception shortly after BRCA1/2 test result disclosure. <i>Genetics in Medicine</i> , 2011, 13, 800-811.	1.1	25
112	Allele-specific regulation of FGFR2 expression is cell type-dependent and may increase breast cancer risk through a paracrine stimulus involving FGF10. <i>Breast Cancer Research</i> , 2011, 13, R72.	2.2	35
113	The MDM2 Promoter SNP285C/309G Haplotype Diminishes Sp1 Transcription Factor Binding and Reduces Risk for Breast and Ovarian Cancer in Caucasians. <i>Cancer Cell</i> , 2011, 19, 273-282.	7.7	104
114	A whisper-game perspective on the family communication of DNA-test results: a retrospective study on the communication process of BRCA1/2-test results between proband and relatives. <i>Familial Cancer</i> , 2011, 10, 87-96.	0.9	48
115	Body weight and risk of breast cancer in BRCA1/2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2011, 126, 193-202.	1.1	59
116	Haplotype structure in Ashkenazi Jewish BRCA1 and BRCA2 mutation carriers. <i>Human Genetics</i> , 2011, 130, 685-699.	1.8	18
117	Genetic testing for familial/hereditary breast cancer: comparison of guidelines and recommendations from the UK, France, the Netherlands and Germany. <i>Journal of Community Genetics</i> , 2011, 2, 53-69.	0.5	59
118	Family communication matters: The impact of telling relatives about unclassified variants and uninformative DNA-test results. <i>Genetics in Medicine</i> , 2011, 13, 333-341.	1.1	33
119	Common alleles at 6q25.1 and 1p11.2 are associated with breast cancer risk for BRCA1 and BRCA2 mutation carriers. <i>Human Molecular Genetics</i> , 2011, 20, 3304-3321.	1.4	68
120	Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. <i>Human Molecular Genetics</i> , 2011, 20, 4732-4747.	1.4	32
121	Physical activity and the risk of breast cancer in BRCA1/2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2010, 120, 235-244.	1.1	79
122	Common Breast Cancer Susceptibility Alleles and the Risk of Breast Cancer for BRCA1 and BRCA2 Mutation Carriers: Implications for Risk Prediction. <i>Cancer Research</i> , 2010, 70, 9742-9754.	0.4	169
123	Evidence for SMAD3 as a modifier of breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2010, 12, R102.	2.2	25
124	TP53 germline mutation testing in 180 families suspected of Li-Fraumeni syndrome: mutation detection rate and relative frequency of cancers in different familial phenotypes. <i>Journal of Medical Genetics</i> , 2010, 47, 421-428.	1.5	254
125	Disentangling the Babylonian speech confusion in genetic counseling: An analysis of the reliability and validity of the nomenclature for BRCA1/2 DNA-test results other than pathogenic. <i>Genetics in Medicine</i> , 2009, 11, 742-749.	1.1	6
126	A simple method for co-segregation analysis to evaluate the pathogenicity of unclassified variants; BRCA1 and BRCA2 as an example. <i>BMC Cancer</i> , 2009, 9, 211.	1.1	57

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127	Intronic variants in BRCA1 and BRCA2 that affect RNA splicing can be reliably selected by splice-site prediction programs. <i>Human Mutation</i> , 2009, 30, 107-114.	1.1	97
128	Differences and similarities in breast cancer risk assessment models in clinical practice: which model to choose?. <i>Breast Cancer Research and Treatment</i> , 2009, 115, 381-390.	1.1	88
129	A 7 Mb region within 11q13 may contain a high penetrance gene for breast cancer. <i>Breast Cancer Research and Treatment</i> , 2009, 118, 151-159.	1.1	23
130	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009, 41, 585-590.	9.4	434
131	A method to assess the clinical significance of unclassified variants in the BRCA1 and BRCA2 genes based on cancer family history. <i>Breast Cancer Research</i> , 2009, 11, R8.	2.2	45
132	The counsellors' view of an unclassified variant in BRCA1/2: recall, interpretation, and impact on life. <i>Psycho-Oncology</i> , 2008, 17, 822-830.	1.0	112
133	Genome-wide linkage scan in Dutch hereditary non-BRCA1/2 breast cancer families identifies 9q21-q22 as a putative breast cancer susceptibility locus. <i>Genes Chromosomes and Cancer</i> , 2008, 47, 947-956.	1.5	16
134	A family history of breast cancer will not predict female early onset breast cancer in a population-based setting. <i>BMC Cancer</i> , 2008, 8, 203.	1.1	9
135	Design of the BRISC study: a multicentre controlled clinical trial to optimize the communication of breast cancer risks in genetic counselling. <i>BMC Cancer</i> , 2008, 8, 283.	1.1	17
136	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	1.5	315
137	Putting it all behind: long-term psychological impact of an inconclusive DNA test result for breast cancer. <i>Genetics in Medicine</i> , 2008, 10, 745-750.	1.1	16
138	Prediction of BRCA1/2 mutation status in patients with ovarian cancer from a hospital-based cohort. <i>Genetics in Medicine</i> , 2007, 9, 173-179.	1.1	17
139	Prognostic factors for hereditary cancer distress six months after BRCA1/2 or HNPCC genetic susceptibility testing. <i>European Journal of Cancer</i> , 2007, 43, 71-77.	1.3	36
140	Clinical correlates of low-risk variants in FGFR2, TNRC9, MAP3K1, LSP1 and 8q24 in a Dutch cohort of incident breast cancer cases. <i>Breast Cancer Research</i> , 2007, 9, R78.	2.2	64
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