

Marc Tischkowitz

List of Publications by Year in descending order

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Version: 2024-02-01

238
papers

23,379
citations

12597

71
h-index

10679

143
g-index

255
all docs

255
docs citations

255
times ranked

28887
citing authors

#	ARTICLE	IF	CITATIONS
1	UK recommendations for <i>SDHA</i> germline genetic testing and surveillance in clinical practice. <i>Journal of Medical Genetics</i> , 2023, 60, 107-111.	1.5	4
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	Quantifying prediction of pathogenicity for within-codon concordance (PM5) using 7541 functional classifications of <i>BRCA1</i> and <i>MSH2</i> missense variants. <i>Genetics in Medicine</i> , 2022, 24, 552-563.	1.1	5
4	Comprehensive epithelial tubo-ovarian cancer risk prediction model incorporating genetic and epidemiological risk factors. <i>Journal of Medical Genetics</i> , 2022, 59, 632-643.	1.5	33
5	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
6	Quantifying evidence toward pathogenicity for rare phenotypes: The case of succinate dehydrogenase genes, <i>SDHB</i> and <i>SDHD</i> . <i>Genetics in Medicine</i> , 2022, 24, 41-50.	1.1	5
7	Cancer Risks Associated With <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>Journal of Clinical Oncology</i> , 2022, 40, 1529-1541.	0.8	90
8	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
9	Multilocus Inherited Neoplasia Allele Syndrome (MINAS): an update. <i>European Journal of Human Genetics</i> , 2022, 30, 265-270.	1.4	12
10	<i>BRCA1</i> and <i>BRCA2</i> pathogenic variants and prostate cancer risk: systematic review and meta-analysis. <i>British Journal of Cancer</i> , 2022, 126, 1067-1081.	2.9	18
11	Targeted long-read sequencing identifies missing pathogenic variants in unsolved Werner syndrome cases. <i>Journal of Medical Genetics</i> , 2022, 59, 1087-1094.	1.5	14
12	Personalised Risk Prediction in Hereditary Breast and Ovarian Cancer: A Protocol for a Multi-Centre Randomised Controlled Trial. <i>Cancers</i> , 2022, 14, 2716.	1.7	10
13	UKCGG Consensus Group guidelines for the management of patients with constitutional <i>TP53</i> pathogenic variants. <i>Journal of Medical Genetics</i> , 2021, 58, 135-139.	1.5	23
14	Combining evidence for and against pathogenicity for variants in cancer susceptibility genes: CanVIG-UK consensus recommendations. <i>Journal of Medical Genetics</i> , 2021, 58, 297-304.	1.5	28
15	Hereditary Diffuse Gastric Cancer: Approaches to Screening, Surveillance, and Treatment. <i>Annual Review of Medicine</i> , 2021, 72, 263-280.	5.0	15
16	A no-deal Brexit will be detrimental to people with rare diseases. <i>Lancet, The</i> , 2021, 397, 20.	6.3	4
17	Pathogenic germline variants in patients with features of hereditary renal cell carcinoma: Evidence for further locus heterogeneity. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 5-16.	1.5	10
18	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

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19	Update: variable implementation of the 2018 UKCGG/UKGTN guidelines for breast cancer gene panel tests offered by UK genetics services. <i>Journal of Medical Genetics</i> , 2021, 58, 579-580.	1.5	1
20	Quantitative evidence evaluation for singleton rare missense variants in rare distinctive adult-onset phenotypes: the exemplar of SDHB and SDHD. <i>Molecular Genetics and Metabolism</i> , 2021, 132, S37.	0.5	0
21	Management of individuals with germline variants in PALB2: a clinical practice resource of the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2021, 23, 1416-1423.	1.1	34
22	Clonal hematopoiesis and therapy-related myeloid neoplasms following neuroblastoma treatment. <i>Blood</i> , 2021, 137, 2992-2997.	0.6	19
23	A mosaic PIK3CA variant in a young adult with diffuse gastric cancer: case report. <i>European Journal of Human Genetics</i> , 2021, 29, 1354-1358.	1.4	9
24	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 BRCA1 or BRCA2 pathogenic variant. <i>Genetics in Medicine</i> , 2021, 23, 1726-1737.	1.1	16
25	Surveillance recommendations for DICER1 pathogenic variant carriers: a report from the SIOPE Host Genome Working Group and CanGene-CanVar Clinical Guideline Working Group. <i>Familial Cancer</i> , 2021, 20, 337-348.	0.9	19
26	Investigating the clinical, pathological and molecular profile of oncocytic adrenocortical neoplasms: a case series and literature review. <i>Endocrine Oncology</i> , 2021, 1, 33-44.	0.1	0
27	CanRisk Tool—A Web Interface for the Prediction of Breast and Ovarian Cancer Risk and the Likelihood of Carrying Genetic Pathogenic Variants. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 469-473.	1.1	98
28	Overview of hereditary breast and ovarian cancer (HBOC) guidelines across Europe. <i>European Journal of Medical Genetics</i> , 2021, 64, 104350.	0.7	22
29	Assessing BRCA1 activity in DNA damage repair using human induced pluripotent stem cells as an approach to assist classification of BRCA1 variants of uncertain significance. <i>PLoS ONE</i> , 2021, 16, e0260852.	1.1	2
30	A functionally impaired missense variant identified in French Canadian families implicates FANCI as a candidate ovarian cancer-predisposing gene. <i>Genome Medicine</i> , 2021, 13, 186.	3.6	12
31	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
32	A systematic review of predicted pathogenic PALB2 variants: an analysis of mutational overlap between epithelial cancers. <i>Journal of Human Genetics</i> , 2020, 65, 199-205.	1.1	12
33	Prostate Cancer Risks for Male BRCA1 and BRCA2 Mutation Carriers: A Prospective Cohort Study. <i>European Urology</i> , 2020, 77, 24-35.	0.9	124
34	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
35	Cancer Risks Associated With Germline <i>PALB2</i> Pathogenic Variants: An International Study of 524 Families. <i>Journal of Clinical Oncology</i> , 2020, 38, 674-685.	0.8	270
36	Whole Exome Sequencing Identifies Candidate Genes Associated with Hereditary Predisposition to Uveal Melanoma. <i>Ophthalmology</i> , 2020, 127, 668-678.	2.5	27

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37	Sporadic implementation of UK familial mammographic surveillance guidelines 15 years after original publication. <i>British Journal of Cancer</i> , 2020, 122, 329-332.	2.9	4
38	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	1.1	82
39	Hereditary diffuse gastric cancer: updated clinical practice guidelines. <i>Lancet Oncology</i> , The, 2020, 21, e386-e397.	5.1	237
40	Association of germline variation with the survival of women with BRCA1/2 pathogenic variants and breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 44.	2.3	5
41	A case-control study of the joint effect of reproductive factors and radiation treatment for first breast cancer and risk of contralateral breast cancer in the WECARE study. <i>Breast</i> , 2020, 54, 62-69.	0.9	3
42	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581.	9.4	265
43	Cancer Surveillance Guideline for individuals with PTEN hamartoma tumour syndrome. <i>European Journal of Human Genetics</i> , 2020, 28, 1387-1393.	1.4	63
44	Cancer Variant Interpretation Group UK (CanVIG-UK): an exemplar national subspecialty multidisciplinary network. <i>Journal of Medical Genetics</i> , 2020, 57, 829-834.	1.5	30
45	Personalized early detection and prevention of breast cancer: ENVISION consensus statement. <i>Nature Reviews Clinical Oncology</i> , 2020, 17, 687-705.	12.5	178
46	Prostate Cancer Risk by BRCA2 Genomic Regions. <i>European Urology</i> , 2020, 78, 494-497.	0.9	6
47	Evaluating clinician acceptability of the prototype CanRisk tool for predicting risk of breast and ovarian cancer: A multi-methods study. <i>PLoS ONE</i> , 2020, 15, e0229999.	1.1	40
48	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
49	Radiation Treatment, <i>ATM</i> , <i>BRCA1/2</i> , and <i>CHEK2</i> *1100delC Pathogenic Variants and Risk of Contralateral Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1275-1279.	3.0	21
50	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
51	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
52	Alcohol Consumption, Cigarette Smoking, and Risk of Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Results from The BRCA1 and BRCA2 Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 368-378.	1.1	24
53	Characterization of renal cell carcinoma-associated constitutional chromosome abnormalities by genome sequencing. <i>Genes Chromosomes and Cancer</i> , 2020, 59, 333-347.	1.5	10
54	Risk-reducing salpingo-oophorectomy, natural menopause, and breast cancer risk: an international prospective cohort of BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2020, 22, 8.	2.2	41

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55	Small-Cell Carcinoma of the Ovary, Hypercalcemic Type—Genetics, New Treatment Targets, and Current Management Guidelines. <i>Clinical Cancer Research</i> , 2020, 26, 3908-3917.	3.2	82
56	Title is missing!. , 2020, 15, e0229999.		0
57	Title is missing!. , 2020, 15, e0229999.		0
58	Title is missing!. , 2020, 15, e0229999.		0
59	Title is missing!. , 2020, 15, e0229999.		0
60	Genotype, extrapyramidal features, and severity of variant ataxia—telangiectasia. <i>Annals of Neurology</i> , 2019, 85, 170-180.	2.8	58
61	Psychosocial impact of undergoing prostate cancer screening for men with <i>BRCA1</i> or <i>BRCA2</i> mutations. <i>BJU International</i> , 2019, 123, 284-292.	1.3	9
62	Homologous recombination DNA repair defects in PALB2-associated breast cancers. <i>Npj Breast Cancer</i> , 2019, 5, 23.	2.3	39
63	Caveat Emptor: The Perils of Panel Testing in Hereditary Breast Cancer. <i>Journal of Clinical Oncology</i> , 2019, 37, 2176-2177.	0.8	2
64	<i>PTEN</i> Hamartoma tumor syndrome in childhood: A review of the clinical literature. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2019, 181, 591-610.	0.7	46
65	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	2.3	28
66	Malta (MYH9 Associated Elastin Aggregation) Syndrome: Germline Variants in MYH9 Cause Rare Sweat Duct Proliferations and Irregular Elastin Aggregations. <i>Journal of Investigative Dermatology</i> , 2019, 139, 2238-2241.e6.	0.3	5
67	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
68	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. <i>JAMA Network Open</i> , 2019, 2, e1912259.	2.8	5
69	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
70	Large scale multifactorial likelihood quantitative analysis of <i>BRCA1</i> and <i>BRCA2</i> variants: An ENIGMA resource to support clinical variant classification. <i>Human Mutation</i> , 2019, 40, 1557-1578.	1.1	102
71	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
72	Flexible and scalable diagnostic filtering of genomic variants using G2P with Ensembl VEP. <i>Nature Communications</i> , 2019, 10, 2373.	5.8	86

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73	<i>BRCA1</i> and <i>BRCA2</i> pathogenic sequence variants in women of African origin or ancestry. <i>Human Mutation</i> , 2019, 40, 1781-1796.	1.1	26
74	The clinical presentation caused by truncating <i>CHD8</i> variants. <i>Clinical Genetics</i> , 2019, 96, 72-84.	1.0	32
75	Towards controlled terminology for reporting germline cancer susceptibility variants: an ENIGMA report. <i>Journal of Medical Genetics</i> , 2019, 56, 347-357.	1.5	32
76	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
77	The Manchester International Consensus Group recommendations for the management of gynecological cancers in Lynch syndrome. <i>Genetics in Medicine</i> , 2019, 21, 2390-2400.	1.1	153
78	Letter from the New Editor in Chief. <i>Genetical Research</i> , 2019, 101, e2.	0.3	0
79	Clinical implications of germline mutations in breast cancer genes: RECQL. <i>Breast Cancer Research and Treatment</i> , 2019, 174, 553-560.	1.1	14
80	BOADICEA: a comprehensive breast cancer risk prediction model incorporating genetic and nongenetic risk factors. <i>Genetics in Medicine</i> , 2019, 21, 1708-1718.	1.1	415
81	Height and Body Mass Index as Modifiers of Breast Cancer Risk in <i>BRCA1</i> / <i>BRCA2</i> Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019, 111, 350-364.	3.0	30
82	Clinicians' use of breast cancer risk assessment tools according to their perceived importance of breast cancer risk factors: an international survey. <i>Journal of Community Genetics</i> , 2019, 10, 61-71.	0.5	7
83	Penetrance estimates for <i>BRCA1</i> , <i>BRCA2</i> (also applied to Lynch syndrome) based on presymptomatic testing: a new unbiased method to assess risk?. <i>Journal of Medical Genetics</i> , 2018, 55, 442-448.	1.5	1
84	Consensus for genes to be included on cancer panel tests offered by UK genetics services: guidelines of the UK Cancer Genetics Group. <i>Journal of Medical Genetics</i> , 2018, 55, 372-377.	1.5	88
85	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.	1.1	224
86	Evaluation of universal immunohistochemical screening of sebaceous neoplasms in a service setting. <i>Clinical and Experimental Dermatology</i> , 2018, 43, 410-415.	0.6	13
87	Histone Lysine Methylases and Demethylases in the Landscape of Human Developmental Disorders. <i>American Journal of Human Genetics</i> , 2018, 102, 175-187.	2.6	204
88	Germline pathogenic variants in <i>PALB2</i> and other cancer-predisposing genes in families with hereditary diffuse gastric cancer without <i>CDH1</i> mutation: a whole-exome sequencing study. <i>The Lancet Gastroenterology and Hepatology</i> , 2018, 3, 489-498.	3.7	87
89	pedigreejs: a web-based graphical pedigree editor. <i>Bioinformatics</i> , 2018, 34, 1069-1071.	1.8	9
90	Risks of breast or ovarian cancer in <i>BRCA1</i> or <i>BRCA2</i> predictive test negatives: findings from the EMBRACE study. <i>Genetics in Medicine</i> , 2018, 20, 1575-1582.	1.1	15

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91	Use of the BOADICEA Web Application in clinical practice: appraisals by clinicians from various countries. <i>Familial Cancer</i> , 2018, 17, 31-41.	0.9	9
92	Comparative study of endoscopic surveillance in hereditary diffuse gastric cancer according to CDH1 mutation status. <i>Gastrointestinal Endoscopy</i> , 2018, 87, 408-418.	0.5	85
93	Clinical implications of germline mutations in breast cancer: TP53. <i>Breast Cancer Research and Treatment</i> , 2018, 167, 417-423.	1.1	112
94	Breast Cancer Family History and Contralateral Breast Cancer Risk in Young Women: An Update From the Women's Environmental Cancer and Radiation Epidemiology Study. <i>Journal of Clinical Oncology</i> , 2018, 36, 1513-1520.	0.8	44
95	p.Val804Met, the Most Frequent Pathogenic Mutation in RET, Confers a Very Low Lifetime Risk of Medullary Thyroid Cancer. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2018, 103, 4275-4282.	1.8	39
96	De Novo Truncating Mutations in WASF1 Cause Intellectual Disability with Seizures. <i>American Journal of Human Genetics</i> , 2018, 103, 144-153.	2.6	36
97	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2018, 78, 5419-5430.	0.4	54
98	Comprehensive Cancer-Predisposition Gene Testing in an Adult Multiple Primary Tumor Series Shows a Broad Range of Deleterious Variants and Atypical Tumor Phenotypes. <i>American Journal of Human Genetics</i> , 2018, 103, 3-18.	2.6	46
99	Biallelic Mutation of ARHGEF18, Involved in the Determination of Epithelial Apicobasal Polarity, Causes Adult-Onset Retinal Degeneration. <i>American Journal of Human Genetics</i> , 2017, 100, 334-342.	2.6	26
100	Genetics of gynaecological cancers. <i>Best Practice and Research in Clinical Obstetrics and Gynaecology</i> , 2017, 42, 114-124.	1.4	13
101	Prevalence and architecture of de novo mutations in developmental disorders. <i>Nature</i> , 2017, 542, 433-438.	13.7	1,211
102	Pathology update to the Manchester Scoring System based on testing in over 4000 families. <i>Journal of Medical Genetics</i> , 2017, 54, 674-681.	1.5	51
103	Risks of Breast, Ovarian, and Contralateral Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>JAMA - Journal of the American Medical Association</i> , 2017, 317, 2402.	3.8	1,898
104	Universal <i>BRCA1/BRCA2</i> Testing for Ovarian Cancer Patients is Welcomed, but with Care: How Women and Staff Contextualize Experiences of Expanded Access. <i>Journal of Genetic Counseling</i> , 2017, 26, 1280-1291.	0.9	14
105	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
106	Compromised <i>BRCA1</i> - <i>PALB2</i> interaction is associated with breast cancer risk. <i>Oncogene</i> , 2017, 36, 4161-4170.	2.6	71
107	Comprehensive Rare Variant Analysis via Whole-Genome Sequencing to Determine the Molecular Pathology of Inherited Retinal Disease. <i>American Journal of Human Genetics</i> , 2017, 100, 75-90.	2.6	343
108	Phenotypic Characterization of <i>EIF2AK4</i> Mutation Carriers in a Large Cohort of Patients Diagnosed Clinically With Pulmonary Arterial Hypertension. <i>Circulation</i> , 2017, 136, 2022-2033.	1.6	111

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109	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
110	Jejunal atresia, periodic fevers and psoriatic arthropathy in Baraitserâ€“Winter malformation syndrome. <i>Clinical Dysmorphology</i> , 2017, 26, 235-237.	0.1	2
111	When to Consider Riskâ€“Reducing Mastectomy in <i>BRCA1/BRCA2</i> Mutation Carriers with Advanced Stage Ovarian Cancer: a Case Study Illustrating the Genetic Counseling Challenges. <i>Journal of Genetic Counseling</i> , 2017, 26, 1173-1178.	0.9	4
112	The use of panel testing in familial breast and ovarian cancer. <i>Clinical Medicine</i> , 2017, 17, 568-572.	0.8	6
113	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.	1.1	18
114	Hormone receptor status of a first primary breast cancer predicts contralateral breast cancer risk in the WECARE study population. <i>Breast Cancer Research</i> , 2017, 19, 83.	2.2	27
115	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
116	PARTNER: Randomised, phase II/III trial to evaluate the safety and efficacy of the addition of olaparib to platinum-based neoadjuvant chemotherapy in triple negative and/or germline <i>BRCA</i> mutated breast cancer patients.. <i>Journal of Clinical Oncology</i> , 2017, 35, TPS591-TPS591.	0.8	7
117	Germline whole exome sequencing and large-scale replication identifies <i>FANCM</i> as a likely high grade serous ovarian cancer susceptibility gene. <i>Oncotarget</i> , 2017, 8, 50930-50940.	0.8	43
118	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	1.5	174
119	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	2.2	31
120	A comparison of the yield from endoscopic surveillance in detecting early gastric cancer in <i>CDH1</i> +ve versus <i>CDH1</i> ^-ve HDGC families. <i>European Journal of Surgical Oncology</i> , 2016, 42, S246-S247.	0.5	0
121	Incorporating truncating variants in <i>PALB2</i> , <i>CHEK2</i> , and <i>ATM</i> into the BOADICEA breast cancer risk model. <i>Genetics in Medicine</i> , 2016, 18, 1190-1198.	1.1	80
122	An international survey of surveillance schemes for unaffected <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2016, 157, 319-327.	1.1	26
123	Body mass index, weight change, and risk of second primary breast cancer in the WECARE study: influence of estrogen receptor status of the first breast cancer. <i>Cancer Medicine</i> , 2016, 5, 3282-3291.	1.3	22
124	Inheritance of deleterious mutations at both <i>BRCA1</i> and <i>BRCA2</i> in an international sample of 32,295 women. <i>Breast Cancer Research</i> , 2016, 18, 112.	2.2	42
125	New paradigms for <i>BRCA1/BRCA2</i> testing in women with ovarian cancer: results of the Genetic Testing in Epithelial Ovarian Cancer (GTEOC) study. <i>Journal of Medical Genetics</i> , 2016, 53, 655-661.	1.5	57
126	Multilocus Inherited Neoplasia Alleles Syndrome. <i>JAMA Oncology</i> , 2016, 2, 373.	3.4	43

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127	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
128	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv315.	3.0	77
129	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
130	Abstract 134: Mutational landscape of breast cancers from PALB2 germline mutation carriers. , 2016, , .		0
131	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2015, 17, 61.	2.2	26
132	Gene-Panel Sequencing and the Prediction of Breast-Cancer Risk. <i>New England Journal of Medicine</i> , 2015, 372, 2243-2257.	13.9	764
133	A homozygous PMS2 founder mutation with an attenuated constitutional mismatch repair deficiency phenotype. <i>Journal of Medical Genetics</i> , 2015, 52, 348-352.	1.5	30
134	Germline Mutation in <i>BRCA1</i> or <i>BRCA2</i> and Ten-Year Survival for Women Diagnosed with Epithelial Ovarian Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 652-657.	3.2	138
135	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
136	Hereditary diffuse gastric cancer: updated clinical guidelines with an emphasis on germline <i>CDH1</i> mutation carriers. <i>Journal of Medical Genetics</i> , 2015, 52, 361-374.	1.5	479
137	Breast cancer risk in women with PALB2 mutations in different populations. <i>Lancet Oncology</i> , The, 2015, 16, e375-e376.	5.1	9
138	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
139	Malignant Peripheral Nerve Sheath Tumor in Cowden Syndrome. <i>Journal of Neuropathology and Experimental Neurology</i> , 2015, 74, 288-292.	0.9	4
140	Heimler Syndrome Is Caused by Hypomorphic Mutations in the Peroxisome-Biogenesis Genes PEX1 and PEX6. <i>American Journal of Human Genetics</i> , 2015, 97, 535-545.	2.6	103
141	Risk reducing salpingectomy and delayed oophorectomy in high risk women: views of cancer geneticists, genetic counsellors and gynaecological oncologists in the UK. <i>Familial Cancer</i> , 2015, 14, 521-530.	0.9	14
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