Christoph Engel

List of Publications by Year
in descending order

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Version: 2024-02-01

272	26,683	11608	7136
papers	citations	70 h-index	153 g-index

289	289	289	26783
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Intensive Insulin Therapy and Pentastarch Resuscitation in Severe Sepsis. New England Journal of Medicine, 2008, 358, 125-139.	13.9	4,141
2	Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA - Journal of the American Medical Association, 2017, 317, 2402.	3.8	1,898
3	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
4	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34.	2.6	711
5	Germline mutations in breast and ovarian cancer pedigrees establish RAD51C as a human cancer susceptibility gene. Nature Genetics, 2010, 42, 410-414.	9.4	638
6	Revised guidelines for the clinical management of Lynch syndrome (HNPCC): recommendations by a group of European experts. Gut, 2013, 62, 812-823.	6.1	630
7	Epidemiology of sepsis in Germany: results from aÂnational prospective multicenter study. Intensive Care Medicine, 2007, 33, 606-618.	3.9	571
8	Breast Cancer Risk Genes â€" Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439.	13.9	532
9	Pathology of Breast and Ovarian Cancers among BRCA1 and BRCA2 Mutation Carriers: Results from the Consortium of Investigators of Modifiers of BRCA1/2 (CIMBA). Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 134-147.	1.1	513
10	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	9.4	493
11	Association of Type and Location ofBRCA1andBRCA2Mutations With Risk of Breast and Ovarian Cancer. JAMA - Journal of the American Medical Association, 2015, 313, 1347.	3.8	390
12	Cancer risks by gene, age, and gender in 6350 carriers of pathogenic mismatch repair variants: findings from the Prospective Lynch Syndrome Database. Genetics in Medicine, 2020, 22, 15-25.	1.1	365
13	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	9.4	356
14	A locus on 19p13 modifies risk of breast cancer in BRCA1 mutation carriers and is associated with hormone receptorâ€"negative breast cancer in the general population. Nature Genetics, 2010, 42, 885-892.	9.4	309
15	Expanded Extracolonic Tumor Spectrum in MUTYH-Associated Polyposis. Gastroenterology, 2009, 137, 1976-1985.e10.	0.6	295
16	Contralateral Breast Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. Journal of Clinical Oncology, 2009, 27, 5887-5892.	0.8	292
17	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
18	The LIFE-Adult-Study: objectives and design of a population-based cohort study with 10,000 deeply phenotyped adults in Germany. BMC Public Health, 2015, 15, 691.	1.2	287

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19	Sleep quality in the general population: psychometric properties of the Pittsburgh Sleep Quality Index, derived from a German community sample of 9284 people. Sleep Medicine, 2017, 30, 57-63.	0.8	274
20	Cancer Risks Associated With Germline PALB2 Pathogenic Variants: An International Study of 524 Families. Journal of Clinical Oncology, 2020, 38, 674-685.	0.8	270
21	Risks of Less Common Cancers in Proven Mutation Carriers With Lynch Syndrome. Journal of Clinical Oncology, 2012, 30, 4409-4415.	0.8	262
22	Common Breast Cancer-Predisposition Alleles Are Associated with Breast Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. American Journal of Human Genetics, 2008, 82, 937-948.	2.6	257
23	Acute renal failure in patients with severe sepsis and septic shock--a significant independent risk factor for mortality: results from the German Prevalence Study. Nephrology Dialysis Transplantation, 2007, 23, 904-909.	0.4	249
24	Genome-Wide Association Study in BRCA1 Mutation Carriers Identifies Novel Loci Associated with Breast and Ovarian Cancer Risk. PLoS Genetics, 2013, 9, e1003212.	1.5	244
25	Lower Incidence of Colorectal Cancer and Later Age of Disease Onset in 27 Families With Pathogenic MSH6 Germline Mutations Compared With Families With MLH1 or MSH2 Mutations: The German Hereditary Nonpolyposis Colorectal Cancer Consortium. Journal of Clinical Oncology, 2004, 22, 4486-4494.	0.8	228
26	The effect of a novel extracorporeal cytokine hemoadsorption device on IL-6 elimination in septic patients: A randomized controlled trial. PLoS ONE, 2017, 12, e0187015.	1.1	227
27	Mutational spectrum in a worldwide study of 29,700 families withBRCA1orBRCA2mutations. Human Mutation, 2018, 39, 593-620.	1.1	224
28	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	9.4	221
29	RAD51 135Gâ†'C Modifies Breast Cancer Risk among BRCA2 Mutation Carriers: Results from a Combined Analysis of 19 Studies. American Journal of Human Genetics, 2007, 81, 1186-1200.	2.6	217
30	Effect of Sodium Selenite Administration and Procalcitonin-Guided Therapy on Mortality in Patients With Severe Sepsis or Septic Shock. JAMA Internal Medicine, 2016, 176, 1266.	2.6	217
31	Effect of Empirical Treatment With Moxifloxacin and Meropenem vs Meropenem on Sepsis-Related Organ Dysfunction in Patients With Severe Sepsis. JAMA - Journal of the American Medical Association, 2012, 307, 2390.	3.8	201
32	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. American Journal of Human Genetics, 2013, 92, 489-503.	2.6	201
33	Effect of Hydrocortisone on Development of Shock Among Patients With Severe Sepsis. JAMA - Journal of the American Medical Association, 2016, 316, 1775.	3.8	197
34	HNPCC-associated small bowel cancer: Clinical and molecular characteristics. Gastroenterology, 2005, 128, 590-599.	0.6	186
35	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. Nature Genetics, 2018, 50, 968-978.	9.4	184
36	Prevalence ofBRCA1/2germline mutations in 21â€…401 families with breast and ovarian cancer. Journal of Medical Genetics, 2016, 53, 465-471.	1.5	179

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37	Impact of compliance with infection management guidelines on outcome in patients with severe sepsis: a prospective observational multi-center study. Critical Care, 2014, 18, R42.	2.5	171
38	Common Breast Cancer Susceptibility Alleles and the Risk of Breast Cancer for BRCA1 and BRCA2 Mutation Carriers: Implications for Risk Prediction. Cancer Research, 2010, 70, 9742-9754.	0.4	169
39	The role of HPV RNA transcription, immune responseâ€related gene expression and disruptive TP53 mutations in diagnostic and prognostic profiling of head and neck cancer. International Journal of Cancer, 2015, 137, 2846-2857.	2.3	169
40	Efficacy of Annual Colonoscopic Surveillance in Individuals With Hereditary Nonpolyposis Colorectal Cancer. Clinical Gastroenterology and Hepatology, 2010, 8, 174-182.	2.4	160
41	Practice and perceptionâ€"A nationwide survey of therapy habits in sepsis*. Critical Care Medicine, 2008, 36, 2719-2725.	0.4	157
42	Prediction of Breast and Prostate Cancer Risks in Male BRCA1 and BRCA2 Mutation Carriers Using Polygenic Risk Scores. Journal of Clinical Oncology, 2017, 35, 2240-2250.	0.8	152
43	Genotype-Phenotype Comparison of German MLH1 and MSH2 Mutation Carriers Clinically Affected With Lynch Syndrome: A Report by the German HNPCC Consortium. Journal of Clinical Oncology, 2006, 24, 4285-4292.	0.8	149
44	Cancer Risks for PMS2-Associated Lynch Syndrome. Journal of Clinical Oncology, 2018, 36, 2961-2968.	0.8	147
45	Effect of a multifaceted educational intervention for anti-infectious measures on sepsis mortality: a cluster randomized trial. Intensive Care Medicine, 2017, 43, 1602-1612.	3.9	143
46	Current practice in nutritional support and its association with mortality in septic patientsâ€"Results from a national, prospective, multicenter study*. Critical Care Medicine, 2008, 36, 1762-1767.	0.4	132
47	Gene panel testing of 5589 <scp>BRCA</scp>1/2â€negative index patients with breast cancer in a routine diagnostic setting: results of the German Consortium for Hereditary Breast and Ovarian Cancer. Cancer Medicine, 2018, 7, 1349-1358.	1.3	126
48	Patients' Perception of Postoperative Pain Management: Validation of the International Pain Outcomes (IPO) Questionnaire. Journal of Pain, 2013, 14, 1361-1370.	0.7	124
49	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
50	Spectrum and frequencies of mutations inMSH2 andMLH1 identified in 1,721 German families suspected of hereditary nonpolyposis colorectal cancer. International Journal of Cancer, 2005, 116, 692-702.	2.3	113
51	The risk of contralateral breast cancer in patients from BRCA1/2 negative high risk families as compared to patients from BRCA1 or BRCA2 positive families: a retrospective cohort study. Breast Cancer Research, 2012, 14, R156.	2.2	112
52	No Difference in Colorectal Cancer Incidence or Stage at Detection by Colonoscopy Among 3 Countries With Different Lynch Syndrome Surveillance Policies. Gastroenterology, 2018, 155, 1400-1409.e2.	0.6	112
53	Ovarian and Breast Cancer Risks Associated With Pathogenic Variants in RAD51C and RAD51D. Journal of the National Cancer Institute, 2020, 112, 1242-1250.	3.0	106
54	A Classification Model for BRCA2 DNA Binding Domain Missense Variants Based on Homology-Directed Repair Activity. Cancer Research, 2013, 73, 265-275.	0.4	103

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55	Large scale multifactorial likelihood quantitative analysis of BRCA1 and BRCA2 variants: An ENIGMA resource to support clinical variant classification. Human Mutation, 2019, 40, 1557-1578.	1.1	102
56	19p13.1 Is a Triple-Negativeâ€“Specific Breast Cancer Susceptibility Locus. Cancer Research, 2012, 72, 1795-1803.	0.4	100
57	Common variants in LSP1, 2q35 and 8q24 and breast cancer risk for BRCA1 and BRCA2 mutation carriers. Human Molecular Genetics, 2009, 18, 4442-4456.	1.4	99
58	Novel strategy for optimal sequential application of clinical criteria, immunohistochemistry and microsatellite analysis in the diagnosis of hereditary nonpolyposis colorectal cancer. International Journal of Cancer, 2006, 118, 115-122.	2.3	98
59	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. American Journal of Human Genetics, 2013, 93, 1046-1060.	2.6	98
60	BRCA1 R1699Q variant displaying ambiguous functional abrogation confers intermediate breast and ovarian cancer risk. Journal of Medical Genetics, 2012, 49, 525-532.	1.5	97
61	Refined histopathological predictors of BRCA1 and BRCA2mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. Breast Cancer Research, 2014, 16, 3419.	2.2	97
62	Validation of the German version of the Mediterranean Diet Adherence Screener (MEDAS) questionnaire. BMC Cancer, 2017, 17, 341.	1.1	95
63	High-risk breast cancer surveillance with MRI: 10-year experience from the German consortium for hereditary breast and ovarian cancer. Breast Cancer Research and Treatment, 2019, 175, 217-228.	1.1	94
64	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. Nature Communications, 2019, 10, 1741.	5.8	90
65	Cancer Risks Associated With BRCA1 and BRCA2 Pathogenic Variants. Journal of Clinical Oncology, 2022, 40, 1529-1541.	0.8	90
66	Evaluating the performance of the breast cancer genetic risk models BOADICEA, IBIS, BRCAPRO and Claus for predictingBRCA1/2mutation carrier probabilities: a study based on 7352 families from the German Hereditary Breast and Ovarian Cancer Consortium. Journal of Medical Genetics, 2013, 50, 360-367.	1.5	88
67	Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. Breast Cancer Research, 2016, 18, 15.	2.2	88
68	Prevalence of pathogenic BRCA1/2 germline mutations among 802 women with unilateral triple-negative breast cancer without family cancer history. BMC Cancer, 2018, 18, 265.	1.1	84
69	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. Genetics in Medicine, 2020, 22, 1653-1666.	1.1	82
70	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
71	Performance of in silico prediction tools for the classification of rare BRCA1/2 missense variants in clinical diagnostics. BMC Medical Genomics, 2018, 11, 35.	0.7	78
72	BRIP1 loss-of-function mutations confer high risk for familial ovarian cancer, but not familial breast cancer. Breast Cancer Research, 2018, 20, 7.	2.2	78

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73	Association Between TAS2R38 Gene Polymorphisms and Colorectal Cancer Risk: A Case-Control Study in Two Independent Populations of Caucasian Origin. PLoS ONE, 2011, 6, e20464.	1.1	77
74	Association Between Loss-of-Function Mutations Within the FANCM Gene and Early-Onset Familial Breast Cancer. JAMA Oncology, 2017, 3, 1245.	3.4	74
75	Common breast cancer susceptibility alleles are associated with tumour subtypes in BRCA1 and BRCA2 mutation carriers: results from the Consortium of Investigators of Modifiers of BRCA1/2. Breast Cancer Research, 2011, 13, R110.	2.2	71
76	Eating Behaviour in the General Population: An Analysis of the Factor Structure of the German Version of the Three-Factor-Eating-Questionnaire (TFEQ) and Its Association with the Body Mass Index. PLoS ONE, 2015, 10, e0133977.	1.1	69
77	Genomic and transcriptomic heterogeneity of colorectal tumours arising in Lynch syndrome. Journal of Pathology, 2017, 243, 242-254.	2.1	69
78	Common alleles at 6q25.1 and 1p11.2 are associated with breast cancer risk for BRCA1 and BRCA2 mutation carriers. Human Molecular Genetics, 2011, 20, 3304-3321.	1.4	68
79	Hereditary Nonpolyposis Colorectal Cancer (HNPCC)/Lynch Syndrome. Deutsches Ärzteblatt International, 2013, 110, 32-8.	0.6	67
80	Reference intervals for leukocyte subsets in adults: Results from a populationâ€based study using 10â€color flow cytometry. Cytometry Part B - Clinical Cytometry, 2015, 88, 270-281.	0.7	65
81	Activating ERBB2/HER2 mutations indicate susceptibility to pan-HER inhibitors in Lynch and Lynch-like colorectal cancer. Gut, 2016, 65, 1296-1305.	6.1	65
82	Systemic and Ocular Determinants of Peripapillary Retinal Nerve Fiber Layer Thickness Measurements in the European Eye Epidemiology (E3) Population. Ophthalmology, 2018, 125, 1526-1536.	2.5	62
83	Common variants associated with breast cancer in genome-wide association studies are modifiers of breast cancer risk in BRCA1 and BRCA2 mutation carriers. Human Molecular Genetics, 2010, 19, 2886-2897.	1.4	60
84	Associations of Pathogenic Variants in MLH1, MSH2, and MSH6 With Risk of Colorectal Adenomas and Tumors and With Somatic Mutations in Patients With Lynch Syndrome. Gastroenterology, 2020, 158, 1326-1333.	0.6	60
85	Automatic Control of Pressure Support for Ventilator Weaning in Surgical Intensive Care Patients. American Journal of Respiratory and Critical Care Medicine, 2012, 185, 637-644.	2.5	59
86	Germline Mutations in Triple-Negative Breast Cancer. Breast Care, 2017, 12, 15-19.	0.8	59
87	The association between unemployment and depressionâ€"Results from the population-based LIFE-adult-study. Journal of Affective Disorders, 2018, 235, 399-406.	2.0	59
88	Variation in the risk of colorectal cancer in families with Lynch syndrome: a retrospective cohort study. Lancet Oncology, The, 2021, 22, 1014-1022.	5.1	58
89	Associations of common breast cancer susceptibility alleles with risk of breast cancer subtypes in BRCA1 and BRCA2 mutation carriers. Breast Cancer Research, 2014, 16, 3416.	2.2	57
90	The â€œunnaturalâ€•history of colorectal cancer in Lynch syndrome: Lessons from colonoscopy surveillance. International Journal of Cancer, 2021, 148, 800-811.	2.3	55

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91	<scp>DNA</scp> methylation array analyses identified breast cancerâ€associated <scp>HYAL2</scp> methylation in peripheral blood. International Journal of Cancer, 2015, 136, 1845-1855.	2.3	53
92	Genome-wide association study of germline variants and breast cancer-specific mortality. British Journal of Cancer, 2019, 120, 647-657.	2.9	52
93	Validation of a brief step-test protocol for estimation of peak oxygen uptake. European Journal of Preventive Cardiology, 2015, 22, 503-512.	0.8	50
94	The BRCA1 c. 5096G>A p.Arg1699Gln (R1699Q) intermediate risk variant: breast and ovarian cancer risk estimation and recommendations for clinical management from the ENIGMA consortium. Journal of Medical Genetics, 2018, 55, 15-20.	1.5	50
95	Development of the Gastrointestinal Dysfunction Score (GIDS) for critically ill patients â€“ A prospective multicenter observational study (iSOFA study). Clinical Nutrition, 2021, 40, 4932-4940.	2.3	49
96	Characterization of the Cancer Spectrum in Men With GermlineBRCA1andBRCA2Pathogenic Variants. JAMA Oncology, 2020, 6, 1218.	3.4	48
97	A computational model of human granulopoiesis to simulate the hematotoxic effects of multicycle polychemotherapy. Blood, 2004, 104, 2323-2331.	0.6	47
98	Common Variants at the 19p13.1 and ZNF365 Loci Are Associated with ER Subtypes of Breast Cancer and Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 645-657.	1.1	47
99	DNA Glycosylases Involved in Base Excision Repair May Be Associated with Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. PLoS Genetics, 2014, 10, e1004256.	1.5	47
100	Breast Cancer Risks and Risk Prediction Models. Breast Care, 2015, 10, 7-12.	0.8	47
101	Arg462Gln sequence variation in the prostate-cancer-susceptibility gene RNASEL and age of onset of hereditary non-polyposis colorectal cancer: a case-control study. Lancet Oncology, The, 2005, 6, 566-572.	5.1	45
102	Double heterozygosity for mutations in BRCA1 and BRCA2 in German breast cancer patients: implications on test strategies and clinical management. Breast Cancer Research and Treatment, 2012, 134, 1229-1239.	1.1	44
103	Germline loss-of-function variants in the BARD1 gene are associated with early-onset familial breast cancer but not ovarian cancer. Breast Cancer Research, 2019, 21, 55.	2.2	44
104	Novel Anthropometry Based on 3D-Bodyscans Applied to a Large Population Based Cohort. PLoS ONE, 2016, 11, e0159887.	1.1	43
105	Prevalence of DSM-5 Mild Neurocognitive Disorder in Dementia-Free Older Adults: Results of the Population-Based LIFE-Adult-Study. American Journal of Geriatric Psychiatry, 2017, 25, 328-339.	0.6	43
106	Sex-Specific Differences in Circumpapillary Retinal Nerve Fiber Layer Thickness. Ophthalmology, 2020, 127, 357-368.	2.5	43
107	Effects of lifestyle intervention in BRCA1/2 mutation carriers on nutrition, BMI, and physical fitness (LIBRE study): study protocol for a randomized controlled trial. Trials, 2016, 17, 368.	0.7	42
108	Lack of association between screening interval and cancer stage in Lynch syndrome may be accounted for by over-diagnosis; a prospective Lynch syndrome database report. Hereditary Cancer in Clinical Practice, 2019, 17, 8.	0.6	42

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109	Symptom Burden and Palliative Care Needs of Patients with Incurable Cancer at Diagnosis and During the Disease Course. Oncologist, 2021, 26, e1058-e1065.	1.9	42
110	The Speaking Voice in the General Population: Normative Data and Associations to Sociodemographic and Lifestyle Factors. Journal of Voice, 2017, 31, 257.e13-257.e24.	0.6	41
111	Risk-reducing salpingo-oophorectomy, natural menopause, and breast cancer risk: an international prospective cohort of BRCA1 and BRCA2 mutation carriers. Breast Cancer Research, 2020, 22, 8.	2.2	41
112	Olfactory function is associated with cognitive performance: results from the population-based LIFE-Adult-Study. Alzheimer's Research and Therapy, 2019, 11, 43.	3.0	40
113	Association of Genomic Domains in BRCA1 and BRCA2 with Prostate Cancer Risk and Aggressiveness. Cancer Research, 2020, 80, 624-638.	0.4	39
114	Association of the VariantsCASP8D302H andCASP10V410I with Breast and Ovarian Cancer Risk inBRCA1andBRCA2Mutation Carriers. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 2859-2868.	1.1	37
115	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. American Journal of Human Genetics, 2015, 97, 22-34.	2.6	37
116	Colorectal surveillance in Lynch syndrome families. Familial Cancer, 2013, 12, 261-265.	0.9	36
117	Absence of association between cyclin D1 (CCND1) G870A polymorphism and age of onset in hereditary nonpolyposis colorectal cancer. Cancer Letters, 2006, 236, 191-197.	3.2	34
118	Association between mental demands at work and cognitive functioning in the general population â€“ results of the health study of the Leipzig research center for civilization diseases (LIFE). Journal of Occupational Medicine and Toxicology, 2014, 9, 23.	0.9	34
119	Assessing Associations between the AURKA-HMMR-TPX2-TUBG1 Functional Module and Breast Cancer Risk in BRCA1/2 Mutation Carriers. PLoS ONE, 2015, 10, e0120020.	1.1	34
120	Oral contraceptive use and ovarian cancer risk for BRCA1/2 mutation carriers: an international cohort study. American Journal of Obstetrics and Gynecology, 2021, 225, 51.e1-51.e17.	0.7	34
121	Oral Contraceptive Use and Breast Cancer Risk: Retrospective and Prospective Analyses From a BRCA1 and BRCA2 Mutation Carrier Cohort Study. JNCI Cancer Spectrum, 2018, 2, pky023.	1.4	33
122	Social factors and the prevalence of social isolation in a population-based adult cohort. Social Psychiatry and Psychiatric Epidemiology, 2022, 57, 1959-1968.	1.6	33
123	Endogenous thrombopoietin serum levels during multicycle chemotherapy. British Journal of Haematology, 1999, 105, 832-838.	1.2	32
124	Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. Human Molecular Genetics, 2011, 20, 4732-4747.	1.4	32
125	Pharmacokinetic and -dynamic modelling of G-CSF derivatives in humans. Theoretical Biology and Medical Modelling, 2012, 9, 32.	2.1	32
126	Breast cancer risk in BRCA1/2 mutation carriers and noncarriers under prospective intensified surveillance. International Journal of Cancer, 2020, 146, 999-1009.	2.3	32

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127	Transcriptomeâ€wide association study of breast cancer risk by estrogenâ€receptor status. Genetic Epidemiology, 2020, 44, 442-468.	0.6	32
128	Cancer risks in Lynch syndrome, Lynch-like syndrome, and familial colorectal cancer type X: a prospective cohort study. BMC Cancer, 2020, 20, 460.	1.1	32
129	Genome-wide analysis associates familial colorectal cancer with increases in copy number variations and a rare structural variation at 12p12.3. Carcinogenesis, 2014, 35, 315-323.	1.3	31
130	Age- and gender-specific norms for the German version of the Three-Factor Eating-Questionnaire (TFEQ). Appetite, 2015, 91, 241-247.	1.8	31
131	Feasibility of structured endurance training and Mediterranean diet in BRCA1 and BRCA2 mutation carriers â€“ an interventional randomized controlled multicenter trial (LIBRE-1). BMC Cancer, 2017, 17, 752.	1.1	31
132	AURKAF31I Polymorphism and Breast Cancer Risk inBRCA1andBRCA2Mutation Carriers: A Consortium of Investigators of Modifiers of BRCA1/2 Study. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 1416-1421.	1.1	30
133	<scp>PAIN OUT</scp>: an international acute pain registry supporting clinicians in decision making and in quality improvement activities. Journal of Evaluation in Clinical Practice, 2014, 20, 1090-1098.	0.9	30
134	Cost-effectiveness of different strategies to prevent breast and ovarian cancer in German women with a BRCA 1 or 2 mutation. European Journal of Health Economics, 2018, 19, 341-353.	1.4	30
135	Height and Body Mass Index as Modifiers of Breast Cancer Risk in BRCA1/2 Mutation Carriers: A Mendelian Randomization Study. Journal of the National Cancer Institute, 2019, 111, 350-364.	3.0	30
136	Validation of the Manchester scoring system for predicting BRCA1/2 mutations in 9,390 families suspected of having hereditary breast and ovarian cancer. International Journal of Cancer, 2014, 135, 2352-2361.	2.3	29
137	Evaluating the performance of clinical criteria for predicting mismatch repair gene mutations in Lynch syndrome: A comprehensive analysis of 3,671 families. International Journal of Cancer, 2014, 135, 69-77.	2.3	28
138	Risk-reducing hysterectomy and bilateral salpingo-oophorectomy in female heterozygotes of pathogenic mismatch repair variants: a Prospective Lynch Syndrome Database report. Genetics in Medicine, 2021, 23, 705-712.	1.1	28
139	Value of upper <scp>gastrointestinal</scp> endoscopy for gastric cancer surveillance in patients with Lynch syndrome. International Journal of Cancer, 2021, 148, 106-114.	2.3	28
140	Age-, sex-, and education-specific norms for an extended CERAD Neuropsychological Assessment Batteryâ€”Results from the population-based LIFE-Adult-Study.. Neuropsychology, 2018, 32, 461-475.	1.0	28
141	A Breast Cancer Risk Haplotype in the Caspase-8 Gene. Cancer Research, 2009, 69, 2724-2728.	0.4	27
142	Age, ocular magnification, and circumpapillary retinal nerve fiber layer thickness. Journal of Biomedical Optics, 2017, 22, 1.	1.4	27
143	Are social conflicts at work associated with depressive symptomatology? Results from the population-based LIFE-Adult-Study. Journal of Occupational Medicine and Toxicology, 2020, 15, 1.	0.9	27
144	Lowâ€risk variants FGFR2, TNRC9 and LSP1 in German familial breast cancer patients. International Journal of Cancer, 2010, 126, 2858-2862.	2.3	26

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145	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. Breast Cancer Research, 2015, 17, 61.	2.2	26
146	Relations between sleep duration with overweight and academic stressâ€"just a matter of the socioeconomic status?. Sleep Health, 2019, 5, 208-215.	1.3	26
147	In vivo effects of interleukin-11 and stem cell factor in combination with erythropoietin in the regulation of erythropoiesis. British Journal of Haematology, 1995, 90, 783-790.	1.2	25
148	Evidence for SMAD3 as a modifier of breast cancer risk in BRCA2mutation carriers. Breast Cancer Research, 2010, 12, R102.	2.2	25
149	The additive effect of p53 Arg72Pro and RNASEL Arg462Gln genotypes on age of disease onset in Lynch syndrome patients with pathogenic germline mutations in MSH2 or MLH1. Cancer Letters, 2007, 252, 55-64.	3.2	24
150	Missense variants in hMLH1 identified in patients from the German HNPCC consortium and functional studies. Familial Cancer, 2011, 10, 273-284.	0.9	24
151	FHL2 expression in peritumoural fibroblasts correlates with lymphatic metastasis in sporadic but not in HNPCC-associated colon cancer. Laboratory Investigation, 2011, 91, 1695-1705.	1.7	24
152	Alcohol Consumption, Cigarette Smoking, and Risk of Breast Cancer for BRCA1 and BRCA2 Mutation Carriers: Results from The BRCA1 and BRCA2 Cohort Consortium. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 368-378.	1.1	24
153	The Human Blood Transcriptome in a Large Population Cohort and Its Relation to Aging and Health. Frontiers in Big Data, 2020, 3, 548873.	1.8	24
154	Polymorphisms of genes coding for ghrelin and its receptor in relation to colorectal cancer risk: a two-step gene-wide case-control study. BMC Gastroenterology, 2010, 10, 112.	0.8	23
155	Targeting of heme oxygenaseâ€1 as a novel immune regulator of neuroblastoma. International Journal of Cancer, 2016, 138, 2030-2042.	2.3	23
156	Targeted Onâ€line SPEâ€LCâ€MS/MS Assay for the Quantitation of 12 Apolipoproteins from Human Blood. Proteomics, 2018, 18, 1700279.	1.3	23
157	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362.	1.4	23
158	Model-based design of chemotherapeutic regimens that account for heterogeneity in leucopoenia. British Journal of Haematology, 2006, 132, 723-735.	1.2	22
159	Identification of aDMBT1polymorphism associated with increased breast cancer risk and decreased promoter activity. Human Mutation, 2010, 31, 60-66.	1.1	22
160	Candidate Genetic Modifiers for Breast and Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 308-316.	1.1	22
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