Volker Arndt

List of Publications by Year in descending order

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243 papers	20,584 citations	11608 70 h-index	12910 131 g-index
268 all docs	268 docs citations	268 times ranked	25738 citing authors

#	Article	IF	Citations
1	Phase III study of the European Organisation for Research and Treatment of Cancer Quality of Life cancer survivorship core questionnaire. Journal of Cancer Survivorship, 2023, 17, 1111-1130.	1.5	6
2	Health and life insurance-related problems in very long-term cancer survivors in Germany: a population-based study. Journal of Cancer Research and Clinical Oncology, 2022, 148, 155-162.	1.2	2
3	Comorbidities, Rather Than Older Age, Are Strongly Associated With Higher Utilization of Healthcare in Colorectal Cancer Survivors. Journal of the National Comprehensive Cancer Network: JNCCN, 2022, 20, 468-478.e7.	2.3	5
4	Rare germline copy number variants (CNVs) and breast cancer risk. Communications Biology, 2022, 5, 65.	2.0	6
5	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. Breast Cancer Research, 2022, 24, 2.	2.2	15
6	Higher Incidence of Diabetes in Cancer Patients Compared to Cancer-Free Population Controls: A Systematic Review and Meta-Analysis. Cancers, 2022, 14, 1808.	1.7	12
7	A Genome-Wide Gene-Based Geneâ€“Environment Interaction Study of Breast Cancer in More than 90,000 Women. Cancer Research Communications, 2022, 2, 211-219.	0.7	6
8	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1077-1089.	1.1	6
9	OUP accepted manuscript. Journal of the National Cancer Institute, 2022, , .	3.0	0
10	Distinct Reproductive Risk Profiles for Intrinsic-Like Breast Cancer Subtypes: Pooled Analysis of Population-Based Studies. Journal of the National Cancer Institute, 2022, 114, 1706-1719.	3.0	14
11	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. Journal of the National Cancer Institute, 2021, 113, 329-337.	3.0	45
12	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. British Journal of Cancer, 2021, 124, 842-854.	2.9	5
13	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. American Journal of Clinical Nutrition, 2021, 113, 1490-1502.	2.2	27
14	Estimation of the Potentially Avoidable Excess Deaths Associated with Socioeconomic Inequalities in Cancer Survival in Germany. Cancers, 2021, 13, 357.	1.7	8
15	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. Nature Communications, 2021, 12, 1078.	5.8	19
16	Genetic architectures of proximal and distal colorectal cancer are partly distinct. Gut, 2021, 70, 1325-1334.	6.1	44
17	Response to Li and Hopper. American Journal of Human Genetics, 2021, 108, 527-529.	2.6	5
18	Trends of incidence, mortality and survival for chronic lymphocytic leukaemia / small lymphocytic lymphoma in Switzerland between 1997 and 2016: a population-based study. Swiss Medical Weekly, 2021, 151, w20463.	0.8	3

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19	Identifying classes of the pain, fatigue, and depression symptom cluster in long-term prostate cancer survivorsâ€"results from the multi-regional Prostate Cancer Survivorship Study in Switzerland (PROCAS). Supportive Care in Cancer, 2021, 29, 6259-6269.	1.0	9
20	Gene-Environment Interactions Relevant to Estrogen and Risk of Breast Cancer: Can Gene-Environment Interactions Be Detected Only among Candidate SNPs from Genome-Wide Association Studies?. Cancers, 2021, 13, 2370.	1.7	4
21	Health-Related Quality of Life in Very Long-Term Cancer Survivors 14â€"24 Years Post-Diagnosis Compared to Population Controls: A Population-Based Study. Cancers, 2021, 13, 2754.	1.7	10
22	Inpatient rehabilitation therapy among colorectal cancer patients â€" utilization and association with prognosis: a cohort study. Acta OncolÃ³gica, 2021, 60, 1000-1010.	0.8	4
23	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. American Journal of Human Genetics, 2021, 108, 1190-1203.	2.6	6
24	Prevalence of benefit finding and posttraumatic growth in long-term cancer survivors: results from a multi-regional population-based survey in Germany. British Journal of Cancer, 2021, 125, 877-883.	2.9	15
25	Distress mediates the relationship between cognitive appraisal of medical care and benefit finding/posttraumatic growth in longâ€term cancer survivors. Cancer, 2021, 127, 3680-3690.	2.0	3
26	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. Breast Cancer Research, 2021, 23, 86.	2.2	7
27	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. British Journal of Cancer, 2021, 125, 1135-1145.	2.9	9
28	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 623-642.	1.1	19
29	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. Scientific Reports, 2021, 11, 19787.	1.6	2
30	Potential to Improve Therapy of Chronic Myeloid Leukemia (CML), Especially for Patients with Older Age: Incidence, Mortality, and Survival Rates of Patients with CML in Switzerland from 1995 to 2017. Cancers, 2021, 13, 6269.	1.7	5
31	The COVID-19 Pandemic and Cancer Patients in Germany: Impact on Treatment, Follow-Up Care and Psychological Burden. Frontiers in Public Health, 2021, 9, 788598.	1.3	14
32	Quality of life, distress, and posttraumatic growth 5Âyears after colorectal cancer diagnosis according to history of inpatient rehabilitation. Journal of Cancer Research and Clinical Oncology, 2021, , 1.	1.2	3
33	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
34	Data from Population-based Cancer Registration for Secondary Data Analysis: Methodological Challenges and Perspectives. Gesundheitswesen, 2020, 82, S62-S71.	0.8	8
35	Age-specific health-related quality of life in disease-free long-term prostate cancer survivors versus male population controlsâ€"results from a population-based study. Supportive Care in Cancer, 2020, 28, 2875-2885.	1.0	9
36	Ageâ€specific prevalence and determinants of depression in longâ€term breast cancer survivors compared to female population controls. Cancer Medicine, 2020, 9, 8713-8721.	1.3	23

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37	Prevalence and severity of longâ€term physical, emotional, and cognitive fatigue across 15 different cancer entities. Cancer Medicine, 2020, 9, 8053-8061.	1.3	33
38	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	2.6	124
39	The relationship between posttraumatic growth and health-related quality of life in adult cancer survivors: A systematic review. Journal of Affective Disorders, 2020, 276, 159-168.	2.0	46
40	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. Nature Genetics, 2020, 52, 572-581.	9.4	265
41	Physical activity and long-term fatigue among colorectal cancer survivors â€“ a population-based prospective study. BMC Cancer, 2020, 20, 438.	1.1	9
42	Healthâ€related quality of life in longâ€term prostate cancer survivors after nerveâ€sparing and nonâ€nerveâ€sparing radical prostatectomyâ€”Results from the multiregional PROCAS study. Cancer Medicine, 2020, 9, 5416-5424.	1.3	6
43	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. Scientific Reports, 2020, 10, 9688.	1.6	2
44	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 860-870.	1.1	26
45	Transcriptomeâ€wide association study of breast cancer risk by estrogenâ€receptor status. Genetic Epidemiology, 2020, 44, 442-468.	0.6	32
46	The relative risk of second primary cancers in Switzerland: a population-based retrospective cohort study. BMC Cancer, 2020, 20, 51.	1.1	39
47	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. Nature Communications, 2020, 11, 312.	5.8	30
48	Association of laparoscopic colectomy versus open colectomy on the long-term health-related quality of life of colon cancer survivors. Surgical Endoscopy and Other Interventional Techniques, 2020, 34, 5593-5603.	1.3	5
49	Cancer-Related Fatigue: Causes and Current Treatment Options. Current Treatment Options in Oncology, 2020, 21, 17.	1.3	174
50	Physical Activity and Long-term Quality of Life among Colorectal Cancer Survivorsâ€”A Population-based Prospective Study. Cancer Prevention Research, 2020, 13, 611-622.	0.7	5
51	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	3.0	129
52	The role of psychosocial resources for long-term breast, colorectal, and prostate cancer survivors: prevalence and associations with health-related quality of life. Supportive Care in Cancer, 2019, 27, 275-286.	1.0	7
53	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. Npj Breast Cancer, 2019, 5, 38.	2.3	28
54	World Health Organization cardiovascular disease risk charts: revised models to estimate risk in 21 global regions. The Lancet Global Health, 2019, 7, e1332-e1345.	2.9	554

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55	Two truncating variants in FANCC and breast cancer risk. Scientific Reports, 2019, 9, 12524.	1.6	5
56	Incidence, mortality, and survival trends of soft tissue and bone sarcoma in Switzerland between 1996 and 2015. Cancer Epidemiology, 2019, 63, 101596.	0.8	43
57	Trends of incidence and survival of patients with chronic myelomonocytic leukemia between 1999 and 2014: A comparison between Swiss and American population-based cancer registries. Cancer Epidemiology, 2019, 59, 51-57.	0.8	14
58	Healthâ€related quality of life in longâ€term survivors with localised prostate cancer by therapyâ€”Results from a populationâ€based study. European Journal of Cancer Care, 2019, 28, e13076.	0.7	19
59	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. Nature Communications, 2019, 10, 1741.	5.8	90
60	The association of cancerâ€related fatigue with allâ€cause mortality of colorectal and endometrial cancer survivors: Results from the populationâ€based PROFILES registry. Cancer Medicine, 2019, 8, 3227-3236.	1.3	22
61	Health-related quality of life in long-term disease-free breast cancer survivors versus female population controls in Germany. Breast Cancer Research and Treatment, 2019, 175, 499-510.	1.1	40
62	Genome-wide association study of germline variants and breast cancer-specific mortality. British Journal of Cancer, 2019, 120, 647-657.	2.9	52
63	Age-specific health-related quality of life in long-term and very long-term colorectal cancer survivors versus population controls â€“ a population-based study. Acta OncolÃ³gica, 2019, 58, 801-810.	0.8	26
64	Return to work after cancer. A multi-regional population-based study from Germany. Acta OncolÃ³gica, 2019, 58, 811-818.	0.8	57
65	Age at Diagnosis and Sex Are Associated With Long-term Deficits in Disease-Specific Health-Related Quality of Life of Survivors of Colon and Rectal Cancer: A Population-Based Study. Diseases of the Colon and Rectum, 2019, 62, 1294-1304.	0.7	15
66	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34.	2.6	711
67	Equalization of four cardiovascular risk algorithms after systematic recalibration: individual-participant meta-analysis of 86 prospective studies. European Heart Journal, 2019, 40, 621-631.	1.0	97
68	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. International Journal of Epidemiology, 2019, 48, 767-780.	0.9	35
69	Cardiovascular Risk Factors Associated With Venous Thromboembolism. JAMA Cardiology, 2019, 4, 163.	3.0	187
70	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377
71	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. International Journal of Epidemiology, 2019, 48, 795-806.	0.9	81
72	The BRCA2 c.68-7TÂ>ÂA variant is not pathogenic: A model for clinical calibration of spliceogenicity. Human Mutation, 2018, 39, 729-741.	1.1	19

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73	Population-based cancer survivorship research: Experiences from Germany and the Netherlands. Journal of Cancer Policy, 2018, 15, 87-91.	0.6	9
74	Risk thresholds for alcohol consumption: combined analysis of individual-participant data for 599â€˜912 current drinkers in 83 prospective studies. Lancet, The, 2018, 391, 1513-1523.	6.3	858
75	Trends of incidence, mortality, and survival of multiple myeloma in Switzerland between 1994 and 2013. Cancer Epidemiology, 2018, 53, 105-110.	0.8	21
76	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. International Journal of Epidemiology, 2018, 47, 526-536.	0.9	88
77	Improvement of relative survival in elderly patients with acute myeloid leukaemia emerging from population-based cancer registries in Switzerland between 2001 and 2013. Cancer Epidemiology, 2018, 52, 55-62.	0.8	8
78	â€œStill a Cancer Patientâ€â€”Associations of Cancer Identity With Patient-Reported Outcomes and Health Care Use Among Cancer Survivors. JNCI Cancer Spectrum, 2018, 2, pky031.	1.4	20
79	Mendelian randomisation study of age at menarche and age at menopause and the risk of colorectal cancer. British Journal of Cancer, 2018, 118, 1639-1647.	2.9	16
80	Incidence Trends of Cervical Cancer and Its Precancerous Lesions in Women of Central Switzerland from 2000 until 2014. Frontiers in Medicine, 2018, 5, 58.	1.2	8
81	Health-related quality of life among long-term (â‰¥5Âyears) prostate cancer survivors by primary intervention: a systematic review. Health and Quality of Life Outcomes, 2018, 16, 22.	1.0	24
82	Potential determinants of physical inactivity among long-term colorectal cancer survivors. Journal of Cancer Survivorship, 2018, 12, 679-690.	1.5	10
83	Quality of life and physical activity in long-term (â‰¥5Âyears post-diagnosis) colorectal cancer survivors - systematic review. Health and Quality of Life Outcomes, 2018, 16, 112.	1.0	72
84	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
85	Quality of life in long-term and very long-term cancer survivors versus population controls in Germany. Acta OncolÃ³gica, 2017, 56, 190-197.	0.8	114
86	Trends of classification, incidence, mortality, and survival of MDS patients in Switzerland between 2001 and 2012. Cancer Epidemiology, 2017, 46, 85-92.	0.8	27
87	BRCA2 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799.	0.4	75
88	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
89	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
90	Geneâ€“environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. International Journal of Cancer, 2017, 141, 1830-1840.	2.3	20

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91	Height, selected genetic markers and prostate cancer risk: results from the PRACTICAL consortium. British Journal of Cancer, 2017, 117, 734-743.	2.9	7
92	Evaluation of completeness of case ascertainment in Swiss cancer registration. European Journal of Cancer Prevention, 2017, 26, S139-S146.	0.6	30
93	Neue Rubrik Epidemiologie. Onkologe, 2017, 23, 89-89.	0.7	0
94	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. Genetics in Medicine, 2017, 19, 599-603.	1.1	67
95	Body mass index and breast cancer survival: a Mendelian randomization analysis. International Journal of Epidemiology, 2017, 46, 1814-1822.	0.9	45
96	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. Breast Cancer Research, 2017, 19, 119.	2.2	43
97	PHIP- a novel candidate breast cancer susceptibility locus on 6q14.1. Oncotarget, 2017, 8, 102769-102782.	0.8	9
98	Abstract 2763: Health-related quality of life among long-term prostate cancer survivors by primary treatment: A systematic review. , 2017, , .		0
99	Improvement of Relative Survival in Elderly Patients with Acute Myeloid Leukemia Emerging from Population-Based Cancer Registries in Switzerland from 2001-2013. Blood, 2017, 130, 863-863.	0.6	0
100	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. Oncotarget, 2016, 7, 80140-80163.	0.8	31
101	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. PLoS Medicine, 2016, 13, e1002105.	3.9	118
102	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. PLoS ONE, 2016, 11, e0160316.	1.1	12
103	Fineâ€scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. International Journal of Cancer, 2016, 139, 1303-1317.	2.3	51
104	PALB2,CHEK2andATMrare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811.	1.5	174
105	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. Breast Cancer Research, 2016, 18, 64.	2.2	31
106	Genetic predisposition to ductal carcinoma in situ of the breast. Breast Cancer Research, 2016, 18, 22.	2.2	43
107	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. Cancer Causes and Control, 2016, 27, 679-693.	0.8	21
108	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. American Journal of Human Genetics, 2016, 99, 903-911.	2.6	59

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109	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	7.7	157
110	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. Nature Communications, 2016, 7, 11375.	5.8	93
111	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
112	Fear of recurrence in long-term cancer survivorsâ€”Do cancer type, sex, time since diagnosis, and social support matter?. Health Psychology, 2016, 35, 1329-1333.	1.3	79
113	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). Scientific Reports, 2016, 6, 32512.	1.6	19
114	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	5.8	50
115	Age- and Tumor Subtypeâ€“Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. Journal of Clinical Oncology, 2016, 34, 2750-2760.	0.8	152
116	No evidence that protein truncating variants inBRIP1are associated with breast cancer risk: implications for gene panel testing. Journal of Medical Genetics, 2016, 53, 298-309.	1.5	94
117	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386.	9.4	125
118	Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. Human Genetics, 2016, 135, 137-154.	1.8	8
119	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. Gynecologic Oncology, 2016, 141, 386-401.	0.6	18
120	RAD51B in Familial Breast Cancer. PLoS ONE, 2016, 11, e0153788.	1.1	26
121	Trends of Classification, Incidence, Mortality, and Survival of MDS Patients in Switzerland Between 2001 and 2012. Blood, 2016, 128, 5539-5539.	0.6	0
122	Investigation of geneâ€environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. International Journal of Cancer, 2015, 136, E685-96.	2.3	34
123	Utilisation of psychosocial and informational services in immigrant and nonâ€immigrant German cancer survivors. Psycho-Oncology, 2015, 24, 919-925.	1.0	19
124	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. Obstetrical and Gynecological Survey, 2015, 70, 758-762.	0.2	0
125	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, .	3.0	428
126	Epidemiology in ovarian carcinoma: Lessons from autopsy. Gynecologic Oncology, 2015, 138, 417-420.	0.6	5

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127	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	1.4	40
128	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. American Journal of Human Genetics, 2015, 96, 5-20.	2.6	76
129	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. Carcinogenesis, 2015, 36, 256-271.	1.3	14
130	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380.	9.4	513
131	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
132	Identification of Novel Genetic Markers of Breast Cancer Survival. Journal of the National Cancer Institute, 2015, 107, .	3.0	56
133	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303.	9.4	357
134	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. Human Molecular Genetics, 2015, 24, 5589-5602.	1.4	67
135	Annexin A1 expression in a pooled breast cancer series: association with tumor subtypes and prognosis. BMC Medicine, 2015, 13, 156.	2.3	51
136	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015, 107, djv219.	3.0	99
137	Fine-Scale Mapping of the 4q24 Locus Identifies Two Independent Loci Associated with Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1680-1691.	1.1	24
138	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. Human Molecular Genetics, 2015, 24, 285-298.	1.4	38
139	Trends in incidence of oesophageal and gastric cancer according to morphology and anatomical location, in Switzerland 1982â€“2011. Swiss Medical Weekly, 2015, 145, w14245.	0.8	8
140	MicroRNA Related Polymorphisms and Breast Cancer Risk. PLoS ONE, 2014, 9, e109973.	1.1	49
141	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. PLoS Genetics, 2014, 10, e1004285.	1.5	39
142	Fear of recurrence in long-term breast cancer survivors-still an issue. Results on prevalence, determinants, and the association with quality of life and depression from the Cancer Survivorship-a multi-regional population-based study. Psycho-Oncology, 2014, 23, 547-554.	1.0	179
143	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2014, 23, 6096-6111.	1.4	53
144	Stageâ€specific associations between beta blocker use and prognosis after colorectal cancer. Cancer, 2014, 120, 1178-1186.	2.0	76

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145	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
146	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46 450 cases and 42 461 controls from the breast cancer association consortium. Human Molecular Genetics, 2014, 23, 1934-1946.	1.4	32
147	Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Geneâ€Environment Interactions. Genetic Epidemiology, 2014, 38, 84-93.	0.6	28
148	FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. British Journal of Cancer, 2014, 110, 1088-1100.	2.9	21
149	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	5.8	105
150	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. Human Molecular Genetics, 2014, 23, 6034-6046.	1.4	12
151	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. Breast Cancer Research, 2014, 16, R51.	2.2	14
152	Fear of recurrence and disease progression in longâ€term (â‰¥5â€‰years) cancer survivorsâ€”a systematic review of quantitative studies. Psycho-Oncology, 2013, 22, 1-11.	1.0	384
153	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
154	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
155	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
156	Circulating 25-hydroxyvitamin D serum concentration and total cancer incidence and mortality: A systematic review and meta-analysis. Preventive Medicine, 2013, 57, 753-764.	1.6	99
157	Vitamin D receptor polymorphism and colorectal cancer-specific and all-cause mortality. Cancer Epidemiology, 2013, 37, 905-907.	0.8	21
158	Genome-wide association studies identify four ER negativeâ€“specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	9.4	374
159	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
160	Quality of life in long-term breast cancer survivors â€“ a 10-year longitudinal population-based study. Acta OncolÃ³gica, 2013, 52, 1119-1128.	0.8	138
161	Evidence of Geneâ€“Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. PLoS Genetics, 2013, 9, e1003284.	1.5	136
162	Vitamin D Receptor Genotype rs731236 (Taq1) and Breast Cancer Prognosis. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 437-442.	1.1	22

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163	19p13.1 Is a Triple-Negativeâ€“Specific Breast Cancer Susceptibility Locus. Cancer Research, 2012, 72, 1795-1803.	0.4	100
164	Absence of Evidence Is Not Evidence of Absence Â– Does Structural Quality Not Matter in Colorectal Cancer PatientsÂ' Satisfaction and Quality of Life?. Onkologie, 2012, 35, 159-160.	1.1	0
165	Genome-wide association analysis identifies three new breast cancer susceptibility loci. Nature Genetics, 2012, 44, 312-318.	9.4	256
166	9q31.2-rs865686 as a Susceptibility Locus for Estrogen Receptor-Positive Breast Cancer: Evidence from the Breast Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1783-1791.	1.1	17
167	11q13 is a susceptibility locus for hormone receptor positive breast cancer. Human Mutation, 2012, 33, 1123-1132.	1.1	35
168	Repeated measures of body mass index and risk of health related outcomes. European Journal of Epidemiology, 2012, 27, 215-224.	2.5	26
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