Peter Andreas Fasching

List of Publications by Year
in descending order

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422 papers	41,100 citations	5261 83 h-index	3181 186 g-index
465 all docs	465 docs citations	465 times ranked	34591 citing authors

#	Article	IF	Citations
1	Pathological complete response and long-term clinical benefit in breast cancer: the CTNeoBC pooled analysis. Lancet, The, 2014, 384, 164-172.	6.3	3,224
2	Definition and Impact of Pathologic Complete Response on Prognosis After Neoadjuvant Chemotherapy in Various Intrinsic Breast Cancer Subtypes. Journal of Clinical Oncology, 2012, 30, 1796-1804.	0.8	2,062
3	Trastuzumab Emtansine for Residual Invasive HER2-Positive Breast Cancer. New England Journal of Medicine, 2019, 380, 617-628.	13.9	1,610
4	Pembrolizumab for Early Triple-Negative Breast Cancer. New England Journal of Medicine, 2020, 382, 810-821.	13.9	1,542
5	Tumour-infiltrating lymphocytes and prognosis in different subtypes of breast cancer: a pooled analysis of 3771 patients treated with neoadjuvant therapy. Lancet Oncology, The, 2018, 19, 40-50.	5.1	1,327
6	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
7	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
8	Neoadjuvant carboplatin in patients with triple-negative and HER2-positive early breast cancer (GeparSixto; GBG 66): a randomised phase 2 trial. Lancet Oncology, The, 2014, 15, 747-756.	5.1	810
9	Association between endometriosis and risk of histological subtypes of ovarian cancer: a pooled analysis of caseâ€“control studies. Lancet Oncology, The, 2012, 13, 385-394.	5.1	753
10	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34.	2.6	711
11	Phase III Randomized Study of Ribociclib and Fulvestrant in Hormone Receptorâ€“Positive, Human Epidermal Growth Factor Receptor 2â€“Negative Advanced Breast Cancer: MONALEESA-3. Journal of Clinical Oncology, 2018, 36, 2465-2472.	0.8	704
12	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. Journal of the National Cancer Institute, 2011, 103, 250-263.	3.0	596
13	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
14	Inherited Mutations in 17 Breast Cancer Susceptibility Genes Among a Large Triple-Negative Breast Cancer Cohort Unselected for Family History of Breast Cancer. Journal of Clinical Oncology, 2015, 33, 304-311.	0.8	521
15	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
16	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
17	Circulating Tumor Cells Predict Survival in Early Average-to-High Risk Breast Cancer Patients. Journal of the National Cancer Institute, 2014, 106, .	3.0	493
18	Neoadjuvant Treatment With Trastuzumab in HER2-Positive Breast Cancer: Results From the GeparQuattro Study. Journal of Clinical Oncology, 2010, 28, 2024-2031.	0.8	487

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19	Overall Survival with Ribociclib plus Fulvestrant in Advanced Breast Cancer. New England Journal of Medicine, 2020, 382, 514-524.	13.9	482
20	Neoadjuvant Chemotherapy and Bevacizumab for HER2-Negative Breast Cancer. New England Journal of Medicine, 2012, 366, 299-309.	13.9	473
21	Association Between CYP2D6 Polymorphisms and Outcomes Among Women With Early Stage Breast Cancer Treated With Tamoxifen. JAMA - Journal of the American Medical Association, 2009, 302, 1429.	3.8	468
22	Pathologic Complete Response After Neoadjuvant Chemotherapy Plus Trastuzumab Predicts Favorable Survival in Human Epidermal Growth Factor Receptor 2â€“Overexpressing Breast Cancer: Results From the TECHNO Trial of the AGO and GBG Study Groups. Journal of Clinical Oncology, 2011, 29, 3351-3357.	0.8	456
23	Event-free Survival with Pembrolizumab in Early Triple-Negative Breast Cancer. New England Journal of Medicine, 2022, 386, 556-567.	13.9	444
24	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, .	3.0	428
25	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	9.4	426
26	Lapatinib versus trastuzumab in combination with neoadjuvant anthracycline-taxane-based chemotherapy (GeparQuinto, GBG 44): a randomised phase 3 trial. Lancet Oncology, The, 2012, 13, 135-144.	5.1	425
27	Genome-wide association studies identify four ER negativeâ€“specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	9.4	374
28	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
29	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	9.4	356
30	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 2013, 45, 362-370.	9.4	326
31	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. Nature Genetics, 2010, 42, 874-879.	9.4	321
32	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
33	Germline Mutation Status, Pathological Complete Response, and Disease-Free Survival in Triple-Negative Breast Cancer. JAMA Oncology, 2017, 3, 1378.	3.4	300
34	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
35	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptorâ€“negative breast cancer. Nature Genetics, 2011, 43, 1210-1214.	9.4	279
36	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. Nature Genetics, 2009, 41, 996-1000.	9.4	276

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37	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
38	Ki67, chemotherapy response, and prognosis in breast cancer patients receiving neoadjuvant treatment. BMC Cancer, 2011, 11, 486.	1.1	260
39	Dose-Response Association of CD8⁺ Tumor-Infiltrating Lymphocytes and Survival Time in High-Grade Serous Ovarian Cancer. JAMA Oncology, 2017, 3, e173290.	3.4	260
40	Genome-wide association analysis identifies three new breast cancer susceptibility loci. Nature Genetics, 2012, 44, 312-318.	9.4	256
41	Clinical and molecular characteristics of HER2-low-positive breast cancer: pooled analysis of individual patient data from four prospective, neoadjuvant clinical trials. Lancet Oncology, The, 2021, 22, 1151-1161.	5.1	248
42	Common variants at 19p13 are associated with susceptibility to ovarian cancer. Nature Genetics, 2010, 42, 880-884.	9.4	235
43	PIK3CA Mutations Are Associated With Lower Rates of Pathologic Complete Response to Antiâ€“Human Epidermal Growth Factor Receptor 2 (HER2) Therapy in Primary HER2-Overexpressing Breast Cancer. Journal of Clinical Oncology, 2014, 32, 3212-3220.	0.8	231
44	Standardized evaluation of tumor-infiltrating lymphocytes in breast cancer: results of the ring studies of the international immuno-oncology biomarker working group. Modern Pathology, 2016, 29, 1155-1164.	2.9	230
45	Impact of treatment characteristics on response of different breast cancer phenotypes: pooled analysis of the German neo-adjuvant chemotherapy trials. Breast Cancer Research and Treatment, 2011, 125, 145-156.	1.1	228
46	Triple-Negative Breast Cancer Risk Genes Identified by Multigene Hereditary Cancer Panel Testing. Journal of the National Cancer Institute, 2018, 110, 855-862.	3.0	225
47	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	9.4	221
48	Efficacy of Margetuximab vs Trastuzumab in Patients With Pretreated ERBB2-Positive Advanced Breast Cancer. JAMA Oncology, 2021, 7, 573.	3.4	217
49	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
50	Capecitabine in Addition to Anthracycline- and Taxane-Based Neoadjuvant Treatment in Patients With Primary Breast Cancer: Phase III GeparQuattro Study. Journal of Clinical Oncology, 2010, 28, 2015-2023.	0.8	194
51	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
52	Identification of nine new susceptibility loci for endometrial cancer. Nature Communications, 2018, 9, 3166.	5.8	178
53	The Contributions of Breast Density and Common Genetic Variation to Breast Cancer Risk. Journal of the National Cancer Institute, 2015, 107, .	3.0	174
54	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. Human Molecular Genetics, 2012, 21, 5373-5384.	1.4	168

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55	CHEK2*1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancerâ€“Specific Death, and Increased Risk of a Second Breast Cancer. Journal of Clinical Oncology, 2012, 30, 4308-4316.	0.8	162
56	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
57	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2011, 20, 3289-3303.	1.4	152
58	Age- and Tumor Subtypeâ€“Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. Journal of Clinical Oncology, 2016, 34, 2750-2760.	0.8	152
59	Neoadjuvant Trastuzumab Emtansine and Pertuzumab in Human Epidermal Growth Factor Receptor 2â€“Positive Breast Cancer: Three-Year Outcomes From the Phase III KRISTINE Study. Journal of Clinical Oncology, 2019, 37, 2206-2216.	0.8	152
60	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. Carcinogenesis, 2014, 35, 1012-1019.	1.3	145
61	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	5.8	144
62	Genome-wide association study identifies a common variant associated with risk of endometrial cancer. Nature Genetics, 2011, 43, 451-454.	9.4	141
63	Evidence of Geneâ€“Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. PLoS Genetics, 2013, 9, e1003284.	1.5	136
64	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
65	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
66	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 1619-1630.	0.9	111
67	Common Breast Cancer Susceptibility Loci Are Associated with Triple-Negative Breast Cancer. Cancer Research, 2011, 71, 6240-6249.	0.4	109
68	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. Nature Communications, 2014, 5, 5303.	5.8	109
69	Quality Assured Health Care in Certified Breast Centers and Improvement of the Prognosis of Breast Cancer Patients. Onkologie, 2011, 34, 362-367.	1.1	106
70	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	5.8	105
71	A Phase II Study of Talazoparib after Platinum or Cytotoxic Nonplatinum Regimens in Patients with Advanced Breast Cancer and Germline BRCA1/2 Mutations (ABRAZO). Clinical Cancer Research, 2019, 25, 2717-2724.	3.2	102
72	Common Breast Cancer Susceptibility Variants in LSP1 and RAD51L1 Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1156-1166.	1.1	101

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73	Risk of Estrogen Receptorâ€“Positive and â€“Negative Breast Cancer and Singleâ€“Nucleotide Polymorphism 2q35-rs13387042. Journal of the National Cancer Institute, 2009, 101, 1012-1018.	3.0	99
74	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015, 107, djv219.	3.0	99
75	Gene panel sequencing in familial breast/ovarian cancer patients identifies multiple novel mutations also in genes others than BRCA1/2. International Journal of Cancer, 2017, 140, 95-102.	2.3	99
76	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
77	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. Nature Communications, 2013, 4, 1627.	5.8	98
78	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
79	The ubiquitin-like molecule interferon-stimulated gene 15 (ISG15) is a potential prognostic marker in human breast cancer. Breast Cancer Research, 2008, 10, R58.	2.2	95
80	Quantification of fatty acid ethyl esters (FAEE) and ethyl glucuronide (EtG) in meconium from newborns for detection of alcohol abuse in a maternal health evaluation study. Analytical and Bioanalytical Chemistry, 2010, 396, 2469-2477.	1.9	95
81	NAB-Paclitaxel Improves Disease-Free Survival in Early Breast Cancer: GBG 69â€“GeparSepto. Journal of Clinical Oncology, 2019, 37, 2226-2234.	0.8	95
82	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
83	The influence of obesity on survival in early, high-risk breast cancer: results from the randomized SUCCESS A trial. Breast Cancer Research, 2015, 17, 129.	2.2	93
84	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. Nature Communications, 2016, 7, 11375.	5.8	93
85	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. Nature Communications, 2019, 10, 1741.	5.8	90
86	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
87	BRCA1/2 Mutations and Bevacizumab in the Neoadjuvant Treatment of Breast Cancer: Response and Prognosis Results in Patients With Triple-Negative Breast Cancer From the GeparQuinto Study. Journal of Clinical Oncology, 2018, 36, 2281-2287.	0.8	86
88	Association of ESR1 gene tagging SNPs with breast cancer risk. Human Molecular Genetics, 2009, 18, 1131-1139.	1.4	84
89	Intense dose-dense epirubicin, paclitaxel, cyclophosphamideÂversus weekly paclitaxel, liposomal doxorubicin (plus carboplatin in triple-negative breast cancer) for neoadjuvant treatment of high-risk early breast cancer (GeparOctoâ€“GBG 84): A randomised phase III trial. European Journal of Cancer, 2019, 106, 181-192.	1.3	84
90	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. Breast Cancer Research, 2010, 12, R110.	2.2	82

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91	Population Distribution of Lifetime Risk of Ovarian Cancer in the United States. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 671-676.	1.1	82
92	The role of genetic breast cancer susceptibility variants as prognostic factors. Human Molecular Genetics, 2012, 21, 3926-3939.	1.4	80
93	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
94	Single nucleotide polymorphisms of the aromatase gene (CYP19A1), HER2/neu status, and prognosis in breast cancer patients. Breast Cancer Research and Treatment, 2008, 112, 89-98.	1.1	77
95	Five endometrial cancer risk loci identified through genome-wide association analysis. Nature Genetics, 2016, 48, 667-674.	9.4	77
96	Association of Pathologic Complete Response with Long-Term Survival Outcomes in Triple-Negative Breast Cancer: A Meta-Analysis. Cancer Research, 2020, 80, 5427-5434.	0.4	77
97	Ki-67 as a prognostic molecular marker in routine clinical use in breast cancer patients. Breast, 2009, 18, 135-141.	0.9	76
98	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
99	A Phase II Randomized Study of Neoadjuvant Letrozole Plus Alpelisib for Hormone Receptor-Positive, Human Epidermal Growth Factor Receptor 2-Negative Breast Cancer (NEO-ORB). Clinical Cancer Research, 2019, 25, 2975-2987.	3.2	76
100	Neoadjuvant chemotherapy with paclitaxel and everolimus in breast cancer patients with non-responsive tumours to epirubicin/cyclophosphamide (EC)Â±bevacizumab â€“ Results of the randomised GeparQuinto study (GBG 44). European Journal of Cancer, 2013, 49, 2284-2293.	1.3	75
101	AGO Recommendations for the Diagnosis and Treatment of Patients with Early Breast Cancer: Update 2019. Breast Care, 2019, 14, 224-245.	0.8	72
102	Influence of mammographic density on the diagnostic accuracy of tumor size assessment and association with breast cancer tumor characteristics. European Journal of Radiology, 2006, 60, 398-404.	1.2	71
103	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 884-895.	0.9	71
104	Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study. Journal of Pathology: Clinical Research, 2018, 4, 250-261.	1.3	70
105	Role of genetic polymorphisms and ovarian cancer susceptibility. Molecular Oncology, 2009, 3, 171-181.	2.1	69
106	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. Human Molecular Genetics, 2015, 24, 5955-5964.	1.4	68
107	Lymphedema in breast cancer survivors: Assessment and information provision in a specialized breast unit. Patient Education and Counseling, 2007, 66, 311-318.	1.0	65
108	Characterizing mammographic images by using generic texture features. Breast Cancer Research, 2012, 14, R59.	2.2	65

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109	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1503-1510.	1.1	64
110	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. Endocrine-Related Cancer, 2016, 23, 77-91.	1.6	62
111	Genetic overlap between endometriosis and endometrial cancer: evidence from crossâ€disease genetic correlation and GWAS metaâ€analyses. Cancer Medicine, 2018, 7, 1978-1987.	1.3	62
112	The 5-HTTLPR polymorphism modulates the influence on environmental stressors on peripartum depression symptoms. Journal of Affective Disorders, 2012, 136, 1192-1197.	2.0	60
113	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
114	Prediction of pathological complete response and prognosis in patients with neoadjuvant treatment for triple-negative breast cancer. BMC Cancer, 2018, 18, 1051.	1.1	59
115	Interdisciplinary Screening, Diagnosis, Therapy and Follow-up of Breast Cancer. Guideline of the DGGG and the DKG (S3-Level, AWMF Registry Number 032/045OL, December 2017) â€“ Part 1 with Recommendations for the Screening, Diagnosis and Therapy of Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2018, 78, 927-948.	0.8	59
116	Five Polymorphisms and Breast Cancer Risk: Results from the Breast Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 1610-1616.	1.1	57
117	Identification of Novel Genetic Markers of Breast Cancer Survival. Journal of the National Cancer Institute, 2015, 107, .	3.0	56
118	BRCA mutations and their influence on pathological complete response and prognosis in a clinical cohort of neoadjuvantly treated breast cancer patients. Breast Cancer Research and Treatment, 2018, 171, 85-94.	1.1	56
119	Impact of disease progression on health-related quality of life in patients with metastatic breast cancer in the PRAEGNANT breast cancer registry. Breast, 2018, 37, 154-160.	0.9	56
120	ABCB1 (MDR1) polymorphisms and ovarian cancer progression and survival: A comprehensive analysis from the Ovarian Cancer Association Consortium and The Cancer Genome Atlas. Gynecologic Oncology, 2013, 131, 8-14.	0.6	55
121	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. Cancer Research, 2015, 75, 2457-2467.	0.4	55
122	Did you drink alcohol during pregnancy? Inaccuracy and discontinuity of women's self-reports: On the way to establish meconium ethyl glucuronide (EtG) as a biomarker for alcohol consumption during pregnancy. Alcohol, 2016, 54, 39-44.	0.8	55
123	Common alleles in candidate susceptibility genes associated with risk and development of epithelial ovarian cancer. International Journal of Cancer, 2011, 128, 2063-2074.	2.3	54
124	Treatment landscape of advanced breast cancer patients with hormone receptor positive HER2 negative tumors â€“ Data from the German PRAEGNANT breast cancer registry. Breast, 2018, 37, 42-51.	0.9	54
125	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. Cancer Research, 2018, 78, 5419-5430.	0.4	54
126	Mammographic density as a risk factor for breast cancer in a German caseâ€“control study. European Journal of Cancer Prevention, 2011, 20, 1-8.	0.6	53

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127	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
128	Genome-wide association study of germline variants and breast cancer-specific mortality. British Journal of Cancer, 2019, 120, 647-657.	2.9	52
129	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
130	AGO Recommendations for the Diagnosis and Treatment of Patients with Early Breast Cancer: Update 2021. Breast Care, 2021, 16, 214-227.	0.8	51
131	Comparison of 6q25 Breast Cancer Hits from Asian and European Genome Wide Association Studies in the Breast Cancer Association Consortium (BCAC). PLoS ONE, 2012, 7, e42380.	1.1	51
132	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. JAMA Oncology, 2022, 8, e216744.	3.4	51
133	Breast Volumetry Using a Three-Dimensional Surface Assessment Technique. Aesthetic Plastic Surgery, 2011, 35, 847-855.	0.5	50
134	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. Human Molecular Genetics, 2015, 24, 1478-1492.	1.4	50
135	MicroRNA Related Polymorphisms and Breast Cancer Risk. PLoS ONE, 2014, 9, e109973.	1.1	49
136	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. Cancer Research, 2019, 79, 505-517.	0.4	49
137	Association Between a Germline OCA2 Polymorphism at Chromosome 15q13.1 and Estrogen Receptorâ€"Negative Breast Cancer Survival. Journal of the National Cancer Institute, 2010, 102, 650-662.	3.0	48
138	Reliability of an e-PRO Tool of EORTC QLQ-C30 for Measurement of Health-Related Quality of Life in Patients With Breast Cancer: Prospective Randomized Trial. Journal of Medical Internet Research, 2017, 19, e322.	2.1	48
139	AGO Recommendations for the Diagnosis and Treatment of Patients with Locally Advanced and Metastatic Breast Cancer: Update 2020. Breast Care, 2020, 15, 294-309.	0.8	47
140	Pooled analysis of the prognostic relevance of progesterone receptor status in five German cohort studies. Breast Cancer Research and Treatment, 2014, 148, 143-151.	1.1	45
141	Association of mammographic density with hormone receptors in invasive breast cancers: Results from a caseâ€only study. International Journal of Cancer, 2012, 131, 2643-2649.	2.3	44
142	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. PLoS ONE, 2015, 10, e0128106.	1.1	44
143	NATALEE: Phase III study of ribociclib (RIBO) + endocrine therapy (ET) as adjuvant treatment in hormone receptorâ€"positive (HR+), human epidermal growth factor receptor 2â€"negative (HER2â€") early breast cancer (EBC).. Journal of Clinical Oncology, 2019, 37, TPS597-TPS597.	0.8	44
144	Genetic variants in the tryptophan hydroxylase 2 gene (TPH2) and depression during and after pregnancy. Journal of Psychiatric Research, 2012, 46, 1109-1117.	1.5	43

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145	Genetic predisposition to ductal carcinoma in situ of the breast. Breast Cancer Research, 2016, 18, 22.	2.2	43
146	Therapy Landscape in Patients with Metastatic HER2-Positive Breast Cancer: Data from the PRAEGNANT Real-World Breast Cancer Registry. Cancers, 2019, 11, 10.	1.7	43
147	Development and Validation of the Gene Expression Predictor of High-grade Serous Ovarian Carcinoma Molecular SubTYPE (PrOTYPE). Clinical Cancer Research, 2020, 26, 5411-5423.	3.2	43
148	Use of complementary and integrative medicine among German breast cancer patients: predictors and implications for patient care within the PRAEGNANT study network. Archives of Gynecology and Obstetrics, 2017, 295, 1239-1245.	0.8	42
149	Association of Germline Variant Status With Therapy Response in High-risk Early-Stage Breast Cancer. JAMA Oncology, 2020, 6, 744.	3.4	42
150	Integrated Analysis of PTEN and p4EBP1 Protein Expression as Predictors for pCR in HER2-Positive Breast Cancer. Clinical Cancer Research, 2016, 22, 2675-2683.	3.2	41
151	Therapy response and prognosis of patients with early breast cancer with low positivity for hormone receptors â€“ An analysis of 2765 patients from neoadjuvant clinical trials. European Journal of Cancer, 2021, 148, 159-170.	1.3	41
152	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	1.4	40
153	Efficacy of neoadjuvant pertuzumab in addition to chemotherapy and trastuzumab in routine clinical treatment of patients with primary breast cancer: a multicentric analysis. Breast Cancer Research and Treatment, 2019, 173, 319-328.	1.1	40
154	Akt and p53 are potential mediators of reduced mammary tumor growth by Chloroquine and the mTOR inhibitor RAD001. Biochemical Pharmacology, 2012, 83, 480-488.	2.0	39
155	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. PLoS Genetics, 2014, 10, e1004285.	1.5	39
156	Socioeconomic status and depression during and after pregnancy in the Franconian Maternal Health Evaluation Studies (FRAMES). Archives of Gynecology and Obstetrics, 2014, 289, 755-763.	0.8	39
157	Patient survival and tumor characteristics associated with CHEK2:p.I157T â€“ findings from the Breast Cancer Association Consortium. Breast Cancer Research, 2016, 18, 98.	2.2	39
158	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. American Journal of Human Genetics, 2020, 107, 837-848.	2.6	39
159	Mutations in BRCA1/2 and Other Panel Genes in Patients With Metastatic Breast Cancer â€”Association With Patient and Disease Characteristics and Effect on Prognosis. Journal of Clinical Oncology, 2021, 39, 1619-1630.	0.8	39
160	Prognostic effect of low-level HER2 expression in patients with clinically negative HER2 status. European Journal of Cancer, 2021, 155, 1-12.	1.3	39
161	Invasive Breast Cancer: Recognition of Molecular Subtypes. Breast Care, 2011, 6, 258-264.	0.8	38
162	Expression of Neuroendocrine Markers in Different Molecular Subtypes of Breast Carcinoma. BioMed Research International, 2014, 2014, 1-9.	0.9	38

#	Article	IF	Citations
163	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
164	Risk, Prediction and Prevention of Hereditary Breast Cancer â€“ Large-Scale Genomic Studies in Times of Big and Smart Data. Geburtshilfe Und Frauenheilkunde, 2018, 78, 481-492.	0.8	38
165	Electronic-Based Patient-Reported Outcomes: Willingness, Needs, and Barriers in Adjuvant and Metastatic Breast Cancer Patients. JMIR Cancer, 2017, 3, e11.	0.9	38
166	Single nucleotide polymorphism D1853N of the ATM gene may alter the risk for breast cancer. Journal of Cancer Research and Clinical Oncology, 2008, 134, 873-882.	1.2	37
167	Early aberrant insulinâ€like growth factor signaling in the progression to endometrial carcinoma is augmented by tamoxifen. International Journal of Cancer, 2008, 123, 2871-2879.	2.3	37
168	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
169	Association of mammographic density with the proliferation marker Ki-67 in a cohort of patients with invasive breast cancer. Breast Cancer Research and Treatment, 2012, 135, 885-892.	1.1	36
170	11q13 is a susceptibility locus for hormone receptor positive breast cancer. Human Mutation, 2012, 33, 1123-1132.	1.1	35
171	Prognostic effect of Ki-67 in common clinical subgroups of patients with HER2-negative, hormone receptor-positive early breast cancer. Breast Cancer Research and Treatment, 2019, 175, 617-625.	1.1	35
172	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. International Journal of Cancer, 2021, 148, 307-319.	2.3	35
173	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
174	Candidate locus analysis of the TERTâ€“CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. Human Genetics, 2015, 134, 231-245.	1.8	34
175	Initial experience with CDK4/6 inhibitor-based therapies compared to antihormone monotherapies in routine clinical use in patients with hormone receptor positive, HER2 negative breast cancer â€" Data from the PRAEGNANT research network for the first 2 years of drug availability in Germany. Breast, 2020, 54, 88-95.	0.9	34
176	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	3.2	33
177	Evaluation of Pathologic Complete Response as a Surrogate for Long-Term Survival Outcomes in Triple-Negative Breast Cancer. Journal of the National Comprehensive Cancer Network: JNCCN, 2020, 18, 1096-1104.	2.3	33
178	Genome-Wide Association Study Identifies a Possible Susceptibility Locus for Endometrial Cancer. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 980-987.	1.1	32
179	TILGen: A Program to Investigate Immune Targets in Breast Cancer Patients - First Results on the Influence of Tumor-Infiltrating Lymphocytes. Breast Care, 2018, 13, 8-14.	0.8	32
180	Outcome after neoadjuvant chemotherapy in estrogen receptor-positive and progesterone receptor-negative breast cancer patients: a pooled analysis of individual patient data from ten prospectively randomized controlled neoadjuvant trials. Breast Cancer Research and Treatment, 2018, 167, 59-71.	1.1	32

#	Article	IF	Citations
181	Mutational Diversity and Therapy Response in Breast Cancer: A Sequencing Analysis in the Neoadjuvant GeparSepto Trial. Clinical Cancer Research, 2019, 25, 3986-3995.	3.2	32
182	Transcriptomeâ€wide association study of breast cancer risk by estrogenâ€receptor status. Genetic Epidemiology, 2020, 44, 442-468.	0.6	32
183	Assessment of Hepatocyte Growth Factor in Ovarian Cancer Mortality. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 1638-1648.	1.1	31
184	HER2 and TOP2A amplification in a hospital-based cohort of breast cancer patients: associations with patient and tumor characteristics. Breast Cancer Research and Treatment, 2014, 145, 193-203.	1.1	31
185	Genetic variants in <scp>VEGF</scp> pathway genes in neoadjuvant breast cancer patients receiving bevacizumab: Results from the randomized phase III <scp>G</scp>epar<scp>Q</scp>uinto study. International Journal of Cancer, 2015, 137, 2981-2988.	2.3	31
186	Efficacy and safety of everolimus plus exemestane in postmenopausal women with hormone receptorâ€positive, human epidermal growth factor receptor 2â€negative locally advanced or metastatic breast cancer: Results of the singleâ€arm, phase IIIB 4EVER trial. International Journal of Cancer, 2019, 144, 877-885.	2.3	31
187	Filtration based assessment of CTCs and CellSearchÂ® based assessment are both powerful predictors of prognosis for metastatic breast cancer patients. BMC Cancer, 2018, 18, 204.	1.1	30
188	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. Nature Communications, 2020, 11, 312.	5.8	30
189	Shared decision-making in metastatic breast cancer: discrepancy between the expected prolongation of life and treatment efficacy between patients and physicians, and influencing factors. Breast Cancer Research and Treatment, 2013, 139, 429-440.	1.1	29
190	A Randomized, Open-label, Presurgical, Window-of-Opportunity Study Comparing the Pharmacodynamic Effects of the Novel Oral SERD AZD9496 with Fulvestrant in Patients with Newly Diagnosed ER+ HER2â^' Primary Breast Cancer. Clinical Cancer Research, 2020, 26, 4242-4249.	3.2	29
191	Germline polymorphisms in an enhancer of PSIP1 are associated with progression-free survival in epithelial ovarian cancer. Oncotarget, 2016, 7, 6353-6368.	0.8	29
192	An Electronic Patient-Reported Outcome Tool for the FACT-B (Functional Assessment of Cancer) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 3 Breast Cancer: Reliability Study. Journal of Medical Internet Research, 2019, 21, e10004.	2.1	29
193	Delivery mode and the course of pre- and postpartum depression. Archives of Gynecology and Obstetrics, 2012, 286, 1407-1412.	0.8	28
194	Hormone replacement therapy and prognosis in ovarian cancer patients. European Journal of Cancer Prevention, 2013, 22, 52-58.	0.6	28
195	Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Geneâ€Environment Interactions. Genetic Epidemiology, 2014, 38, 84-93.	0.6	28
196	Network-Based Integration of GWAS and Gene Expression Identifies a HOX-Centric Network Associated with Serous Ovarian Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1574-1584.	1.1	28
197	Initial Treatment of Patients with Primary Breast Cancer: Evidence, Controversies, Consensus. Geburtshilfe Und Frauenheilkunde, 2017, 77, 633-644.	0.8	28
198	Predicting attention deficit hyperactivity disorder using pregnancy and birth characteristics. Archives of Gynecology and Obstetrics, 2018, 298, 889-895.	0.8	28

#	Article	IF	Citations
199	Diagnosis and Therapy of Triple-Negative Breast Cancer (TNBC) â€“ Recommendations for Daily Routine Practice. Geburtshilfe Und Frauenheilkunde, 2019, 79, 605-617.	0.8	28
200	Confirmation of 5p12 As a Susceptibility Locus for Progesterone-Receptorâ€“Positive, Lower Grade Breast Cancer. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 2222-2231.	1.1	27
201	Meconium Indicators of Maternal Alcohol Abuse during Pregnancy and Association with Patient Characteristics. BioMed Research International, 2014, 2014, 1-11.	0.9	27
202	Computerized patient identification for the EMBRACA clinical trial using real-time data from the PRAEGNANT network for metastatic breast cancer patients. Breast Cancer Research and Treatment, 2016, 158, 59-65.	1.1	27
203	Effects of prenatal alcohol consumption on cognitive development and <scp>ADHD</scp>â€related behaviour in primaryâ€school age: a multilevel study based on meconium ethyl glucuronide. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2018, 59, 110-118.	3.1	27
204	Immune-related Gene Expression Predicts Response to Neoadjuvant Chemotherapy but not Additional Benefit from PD-L1 Inhibition in Women with Early Triple-negative Breast Cancer. Clinical Cancer Research, 2021, 27, 2584-2591.	3.2	27
205	Common germline polymorphisms associated with breast cancer-specific survival. Breast Cancer Research, 2015, 17, 58.	2.2	26
206	ADSCs and adipocytes are the main producers in the autotaxinâ€“lysophosphatidic acid axis of breast cancer and healthy mammary tissue in vitro. BMC Cancer, 2018, 18, 1273.	1.1	26
207	Update Breast Cancer 2019 Part 3 â€“ Current Developments in Early Breast Cancer: Review and Critical Assessment by an International Expert Panel. Geburtshilfe Und Frauenheilkunde, 2019, 79, 470-482.	0.8	26
208	Locoregional recurrence risk after neoadjuvant chemotherapy: A pooled analysis of nine prospective neoadjuvant breast cancer trials. European Journal of Cancer, 2020, 130, 92-101.	1.3	26
209	Assessment of breast cancer tumour size using six different methods. European Radiology, 2011, 21, 1180-1187.	2.3	25
210	Endometriosis as a risk factor for ovarian or endometrial cancer â€” results of a hospital-based caseâ€“control study. BMC Cancer, 2015, 15, 751.	1.1	25
211	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. Endocrine-Related Cancer, 2015, 22, 851-861.	1.6	25
212	Ribociclib plus fulvestrant for advanced breast cancer: Health-related quality-of-life analyses from the MONALEESA-3 study. Breast, 2020, 54, 148-154.	0.9	25
213	International Consensus Conference for Advanced Breast Cancer, Lisbon 2019: ABC5 Consensus â€“ Assessment by a German Group of Experts. Breast Care, 2020, 15, 82-95.	0.8	25
214	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). Journal of Genetics and Genome Research, 2015, 2, .	0.3	25
215	A careful reassessment of anthracycline use in curable breast cancer. Npj Breast Cancer, 2021, 7, 134.	2.3	25
216	Pain perception and detailed visual pain mapping in breast cancer survivors. Breast Cancer Research and Treatment, 2010, 119, 105-110.	1.1	24

#	Article	IF	Citations
217	Two naturally occurring variants of the serotonin receptor geneHTR3Care associated with nausea in pregnancy. Acta Obstetricia Et Gynecologica Scandinavica, 2010, 89, 7-14.	1.3	24
218	Overexpression of SERBP1 (Plasminogen activator inhibitor 1 RNA binding protein) in human breast cancer is correlated with favourable prognosis. BMC Cancer, 2012, 12, 597.	1.1	24
219	CYP2B6*6 is associated with increased breast cancer risk. International Journal of Cancer, 2014, 134, 426-430.	2.3	24
220	Common variants at theCHEK2gene locus and risk of epithelial ovarian cancer. Carcinogenesis, 2015, 36, 1341-1353.	1.3	24
221	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
222	Genetic risk factors for ovarian cancer and their role for endometriosis risk. Gynecologic Oncology, 2017, 145, 142-147.	0.6	24
223	Implementation and Feasibility of Electronic Patient-Reported Outcome (ePRO) Data Entry in the PRAEGNANT Real-Time Advanced and Metastatic Breast Cancer Registry. Geburtshilfe Und Frauenheilkunde, 2017, 77, 870-878.	0.8	24
224	Prediction and clinical utility of a contralateral breast cancer risk model. Breast Cancer Research, 2019, 21, 144.	2.2	24
225	Disseminated tumour cells from the bone marrow of early breast cancer patients: Results from an international pooled analysis. European Journal of Cancer, 2021, 154, 128-137.	1.3	24
226	Prognostic relevance of Ki-67 in the primary tumor for survival after a diagnosis of distant metastasis. Breast Cancer Research and Treatment, 2013, 138, 899-908.	1.1	23
227	Genome-wide association study of subtype-specific epithelial ovarian cancer risk alleles using pooled DNA. Human Genetics, 2014, 133, 481-497.	1.8	23
228	Update Breast Cancer 2018 (Part 2) â€“ Advanced Breast Cancer, Quality of Life and Prevention. Geburtshilfe Und Frauenheilkunde, 2018, 78, 246-259.	0.8	23
229	Polymorphisms in Inflammation Pathway Genes and Endometrial Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 216-223.	1.1	22
230	Epithelialâ€Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. Genetic Epidemiology, 2015, 39, 689-697.	0.6	22
231	Interest in Integrative Medicine Among Postmenopausal Hormone Receptorâ€Positive Breast Cancer Patients in the EvAluate-TM Study. Integrative Cancer Therapies, 2017, 16, 165-175.	0.8	22
232	MyD88 and TLR4 Expression in Epithelial Ovarian Cancer. Mayo Clinic Proceedings, 2018, 93, 307-320.	1.4	22
233	Assessment of mammographic density before and after first full-term pregnancy. European Journal of Cancer Prevention, 2010, 19, 405-412.	0.6	21
234	Genetic Variants in the Genes of the Stress Hormone Signalling Pathway and Depressive Symptoms during and after Pregnancy. BioMed Research International, 2014, 2014, 1-8.	0.9	21

#	Article	IF	Citations
235	Predicting Triple-Negative Breast Cancer Subtype Using Multiple Single Nucleotide Polymorphisms for Breast Cancer Risk and Several Variable Selection Methods. Geburtshilfe Und Frauenheilkunde, 2017, 77, 667-678.	0.8	21
236	Prognostic Impact of Weight Change During Adjuvant Chemotherapy in Patients With High-Risk Early Breast Cancer: Results From the ADEBAR Study. Clinical Breast Cancer, 2018, 18, 175-183.	1.1	21
237	Pathway-Based Analysis of Genome-Wide Association Data Identified SNPs in HMMR as Biomarker for Chemotherapy- Induced Neutropenia in Breast Cancer Patients. Frontiers in Pharmacology, 2018, 9, 158.	1.6	21
238	Update Breast Cancer 2019 Part 2 â€“ Implementation of Novel Diagnostics and Therapeutics in Advanced Breast Cancer Patients in Clinical Practice. Geburtshilfe Und Frauenheilkunde, 2019, 79, 268-280.	0.8	21
239	Human leucocyte antigen class I in hormone receptor-positive, HER2-negative breast cancer: association with response and survival after neoadjuvant chemotherapy. Breast Cancer Research, 2019, 21, 142.	2.2	21
240	Refined cut-off for TP53 immunohistochemistry improves prediction of TP53 mutation status in ovarian mucinous tumors: implications for outcome analyses. Modern Pathology, 2021, 34, 194-206.	2.9	21
241	SNP-SNP interaction analysis of NF-ÎºB signaling pathway on breast cancer survival. Oncotarget, 2015, 6, 37979-37994.	0.8	20
242	Geneâ€“environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. International Journal of Cancer, 2017, 141, 1830-1840.	2.3	20
243	Update Breast Cancer 2018 (Part 1) â€“ Primary Breast Cancer and Biomarkers. Geburtshilfe Und Frauenheilkunde, 2018, 78, 237-245.	0.8	20
244	AGO Recommendations for the Diagnosis and Treatment of Patients with Locally Advanced and Metastatic Breast Cancer: Update 2021. Breast Care, 2021, 16, 228-235.	0.8	20
245	Neoadjuvant giredestrant (GDC-9545) plus palbociclib (P) versus anastrozole (A) plus P in postmenopausal women with estrogen receptorâ€“positive, HER2-negative, untreated early breast cancer (ER+/HER2â€“ eBC): Final analysis of the randomized, open-label, international phase 2 coopERA BC study.. Journal of Clinical Oncology, 2022, 40, 589-589.	0.8	20
246	FemZone trial: a randomized phase II trial comparing neoadjuvant letrozole and zoledronic acid with letrozole in primary breast cancer patients. BMC Cancer, 2014, 14, 66.	1.1	19
247	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. Breast Cancer Research, 2015, 17, 110.	2.2	19
248	Knowledge and attitudes regarding medical research studies among patients with breast cancer and gynecological diseases. BMC Cancer, 2015, 15, 587.	1.1	19
249	Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. Human Genetics, 2016, 135, 741-756.	1.8	19
250	Mammographic density is the main correlate of tumors detected on ultrasound but not on mammography. International Journal of Cancer, 2016, 139, 1967-1974.	2.3	19
251	Children of Prenatally Depressed Mothers: Externalizing and Internalizing Symptoms are Accompanied by Reductions in Specific Social-Emotional Competencies. Journal of Child and Family Studies, 2017, 26, 3135-3144.	0.7	19
252	Update Breast Cancer 2017 â€“ Implementation of Novel Therapies. Geburtshilfe Und Frauenheilkunde, 2017, 77, 1281-1290.	0.8	19

#	Article	IF	Citations
253	Prenatal Alcohol Exposure Is Associated With Adverse Cognitive Effects and Distinct Whole-Genome DNA Methylation Patterns in Primary School Children. Frontiers in Behavioral Neuroscience, 2018, 12, 125.	1.0	19
254	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
255	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
256	Heart Rate Measurement Accuracy of Fitbit Charge 4 and Samsung Galaxy Watch Active2: Device Evaluation Study. JMIR Formative Research, 2022, 6, e33635.	0.7	19
257	Polymorphisms in estrogen metabolism and estrogen pathway genes and the risk of miscarriage. Archives of Gynecology and Obstetrics, 2009, 280, 395-400.	0.8	18
258	Polymorphisms in theRANK/RANKLGenes and Their Effect on Bone Specific Prognosis in Breast Cancer Patients. BioMed Research International, 2014, 2014, 1-7.	0.9	18
259	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. Gynecologic Oncology, 2016, 141, 386-401.	0.6	18
260	Germline variation in ADAMTSL1 is associated with prognosis following breast cancer treatment in young women. Nature Communications, 2017, 8, 1632.	5.8	18
261	Digit ratio (2D:4D) and behavioral symptoms in primary-school aged boys. Early Human Development, 2018, 119, 1-7.	0.8	18
262	Characterization of Molecular Subtypes of Paget Disease of the Breast Using Immunohistochemistry and In Situ Hybridization. Archives of Pathology and Laboratory Medicine, 2019, 143, 206-211.	1.2	18
263	Update Breast Cancer 2019 Part 4 â€“ Diagnostic and Therapeutic Challenges of New, Personalised Therapies for Patients with Early Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2019, 79, 1079-1089.	0.8	18
264	Salivary and hair cortisol as biomarkers of emotional and behavioral symptoms in 6â€“9â€¯year old children. Physiology and Behavior, 2019, 209, 112584.	1.0	18
265	Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. Human Genetics, 2021, 140, 1353-1365.	1.8	18
266	Clinical effectiveness of olaparib monotherapy in germline BRCA-mutated, HER2-negative metastatic breast cancer in a real-world setting: phase IIIb LUCY interim analysis. European Journal of Cancer, 2021, 152, 68-77.	1.3	18
267	Update Breast Cancer 2019 Part 1 â€“ Implementation of Study Results of Novel Study Designs in Clinical Practice in Patients with Early Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2019, 79, 256-267.	0.8	17
268	Anastrozole has an Association between Degree of Estrogen Suppression and Outcomes in Early Breast Cancer and is a Ligand for Estrogen Receptor Î±. Clinical Cancer Research, 2020, 26, 2986-2996.	3.2	17
269	Pooled analysis of two randomized phase III trials (PlanB/SuccessC) comparing six cycles of docetaxel and cyclophosphamide to sequential anthracycline taxane chemotherapy in patients with intermediate and high risk HER2-negative early breast cancer (n=5,923).. Journal of Clinical Oncology, 2018, 36, 522-522.	0.8	17
270	Correlates of the desire for improved cosmetic results after breast-conserving therapy and mastectomy in breast cancer patients. Breast, 2008, 17, 640-645.	0.9	16

#	Article	IF	Citations
271	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. Nature Communications, 2014, 5, 4051.	5.8	16
272	Addition of triple negativity of breast cancer as an indicator for germline mutations in predisposing genes increases sensitivity of clinical selection criteria. BMC Cancer, 2018, 18, 926.	1.1	16
273	Update Breast Cancer 2019 Part 5 â€“ Diagnostic and Therapeutic Challenges of New, Personalised Therapies in Patients with Advanced Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2019, 79, 1090-1099.	0.8	16
274	Development of central nervous system metastases as a first site of metastatic disease in breast cancer patients treated in the neoadjuvant trials GeparQuinto and GeparSixto. Breast Cancer Research, 2019, 21, 60.	2.2	16
275	Translational highlights in breast cancer research and treatment: recent developments with clinical impact. Current Opinion in Obstetrics and Gynecology, 2019, 31, 67-75.	0.9	16
276	Gene Expression Signatures of BRCAness and Tumor Inflammation Define Subgroups of Early-Stage Hormone Receptorâ€“Positive Breast Cancer Patients. Clinical Cancer Research, 2020, 26, 6523-6534.	3.2	16
277	Update Breast Cancer 2020 Part 1 â€“ Early Breast Cancer: Consolidation of Knowledge About Known Therapies. Geburtshilfe Und Frauenheilkunde, 2020, 80, 277-287.	0.8	16
278	Endocrine Treatment for Breast Cancer Patients Revisitedâ€”History, Standard of Care, and Possibilities of Improvement. Cancers, 2021, 13, 5643.	1.7	16
279	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. Gynecologic Oncology, 2015, 136, 542-548.	0.6	15
280	Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. British Journal of Cancer, 2018, 118, 1123-1129.	2.9	15
281	Association between breast cancer risk factors and molecular type in postmenopausal patients with hormone receptor-positive early breast cancer. Breast Cancer Research and Treatment, 2019, 174, 453-461.	1.1	15
282	HLA-G and HLA-F protein isoform expression in breast cancer patients receiving neoadjuvant treatment. Scientific Reports, 2020, 10, 15750.	1.6	15
283	Gemcitabine as adjuvant chemotherapy in patients with high-risk early breast cancerâ€”results from the randomized phase III SUCCESS-A trial. Breast Cancer Research, 2020, 22, 111.	2.2	15
284	The SNP rs6500843 in 16p13.3 is associated with survival specifically among chemotherapy-treated breast cancer patients. Oncotarget, 2015, 6, 7390-7407.	0.8	15
285	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. Breast Cancer Research, 2022, 24, 2.	2.2	15
286	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
287	Association of molecular subtypes with breast cancer risk factors. European Journal of Cancer Prevention, 2015, 24, 484-490.	0.6	14
288	The Clinical Data Intelligence Project. Informatik-Spektrum, 2016, 39, 290-300.	1.0	14

#	Article	IF	Citations
289	Prediction of contralateral breast cancer: external validation of risk calculators in 20 international cohorts. Breast Cancer Research and Treatment, 2020, 181, 423-434.	1.1	14
290	TP53-based interaction analysis identifies cis-eQTL variants for TP53BP2, FBXO28, and FAM53A that associate with survival and treatment outcome in breast cancer. Oncotarget, 2017, 8, 18381-18398.	0.8	14
291	Clinical Relevance of Serum HER2 and Circulating Tumor Cell Detection in Metastatic Breast Cancer Patients. Anticancer Research, 2017, 37, 3117-3128.	0.5	14
292	The impact of anthracyclines in intermediate and high-risk HER2-negative early breast cancerâ€"a pooled analysis of the randomised clinical trials PlanB and SUCCESS C. British Journal of Cancer, 2022, 126, 1715-1724.	2.9	14
293	Inhibition of hyperalgesia by conditioning electrical stimulation in a human pain model. Pain, 2011, 152, 1298-1303.	2.0	13
294	Prognostic molecular markers and neoadjuvant therapy response in anthracycline-treated breast cancer patients. Archives of Gynecology and Obstetrics, 2013, 287, 337-344.	0.8	13
295	Variation in NF-ÎºB Signaling Pathways and Survival in Invasive Epithelial Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1421-1427.	1.1	13
296	Correlation of mammographic density and serum calcium levels in patients with primary breast cancer. Cancer Medicine, 2017, 6, 1473-1481.	1.3	13
297	HLA-J, a Non-Pseudogene as a New Prognostic Marker for Therapy Response and Survival in Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2020, 80, 1123-1133.	0.8	13
298	Mammographic density and prognosis in primary breast cancer patients. Breast, 2021, 59, 51-57.	0.9	13
299	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. Oncotarget, 2016, 7, 72381-72394.	0.8	13
300	Assessment of breast volume changes during human pregnancy using a three-dimensional surface assessment technique in the prospective CGATE study. European Journal of Cancer Prevention, 2014, 23, 151-157.	0.6	12
301	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. Human Molecular Genetics, 2014, 23, 6034-6046.	1.4	12
302	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. PLoS ONE, 2016, 11, e0160316.	1.1	12
303	Neoadjuvant Treatment of Breast Cancer - Advances and Limitations. Breast Care, 2016, 11, 313-314.	0.8	12
304	Semiâ€automated delineation of breast cancer tumors and subsequent materialization using threeâ€dimensional printing (rapid prototyping). Journal of Surgical Oncology, 2017, 115, 238-242.	0.8	12
305	Association between mammographic density and pregnancies relative to age and BMI: a breast cancer case-only analysis. Breast Cancer Research and Treatment, 2017, 166, 701-708.	1.1	12
306	Breast cancer in young women: do BRCA1 or BRCA2 mutations matter?. Lancet Oncology, The, 2018, 19, 150-151.	5.1	12

#	Article	IF	Citations
307	The effect of participation in neoadjuvant clinical trials on outcomes in patients with early breast cancer. Breast Cancer Research and Treatment, 2018, 171, 747-758.	1.1	12
308	Evaluation of soluble carbonic anhydrase IX as predictive marker for efficacy of bevacizumab: A biomarker analysis from the geparquinto phase III neoadjuvant breast cancer trial. International Journal of Cancer, 2019, 145, 857-868.	2.3	12
309	Awareness of breast cancer incidence and risk factors among healthy women in Germany: an update after 10 years. European Journal of Cancer Prevention, 2019, 28, 515-521.	0.6	12
310	Characteristics and Clinical Outcome of Breast Cancer Patients with Asymptomatic Brain Metastases. Cancers, 2020, 12, 2787.	1.7	12
311	Update Breast Cancer 2020 Part 3 â€“ Early Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2020, 80, 1105-1114.	0.8	12
312	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 217-228.	1.1	12
313	Association of genomic variants at the human leukocyte antigen locus with cervical cancer risk, HPV status and gene expression levels. International Journal of Cancer, 2020, 147, 2458-2468.	2.3	12
314	Update Breast Cancer 2020 Part 2 â€“ Advanced Breast Cancer: New Treatments and Implementation of Therapies with Companion Diagnostics. Geburtshilfe Und Frauenheilkunde, 2020, 80, 391-398.	0.8	12
315	Survival analysis of the randomised phase III GeparOcto trial comparing neoadjuvant chemotherapy of intense dose-dense epirubicin, paclitaxel, cyclophosphamide versus weekly paclitaxel, liposomal doxorubicin (plus carboplatin in triple-negative breast cancer) for patients with high-risk early breast cancer. European Journal of Cancer, 2022, 160, 100-111.	1.3	12
316	Pathological Response in the Breast and Axillary Lymph Nodes after Neoadjuvant Systemic Treatment in Patients with Initially Node-Positive Breast Cancer Correlates with Disease Free Survival: An Exploratory Analysis of the GeparOcto Trial. Cancers, 2022, 14, 521.	1.7	12
317	Comprehensive visualization of paresthesia in breast cancer survivors. Archives of Gynecology and Obstetrics, 2014, 290, 135-141.	0.8	11
318	Translational Highlights in Breast and Ovarian Cancer 2019 â€“ Immunotherapy, DNA Repair, PI3K Inhibition and CDK4/6 Therapy. Geburtshilfe Und Frauenheilkunde, 2019, 79, 1309-1319.	0.8	11
319	Update Breast Cancer 2020 Part 4 â€“ Advanced Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2020, 80, 1115-1122.	0.8	11
320	Prevalence of Circulating Tumor Cells After Adjuvant Chemotherapy With or Without Anthracyclines in Patients With HER2-negative, Hormone Receptor-positive Early Breast Cancer. Clinical Breast Cancer, 2017, 17, 279-285.	1.1	10
321	History of Comorbidities and Survival of Ovarian Cancer Patients, Results from the Ovarian Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1470-1473.	1.1	10
322	Updates on the role of receptor activator of nuclear factor ÎºB/receptor activator of nuclear factor ÎºB ligand/osteoprotegerin pathway in breast cancer risk and treatment. Current Opinion in Obstetrics and Gynecology, 2017, 29, 4-11.	0.9	10
323	Initial clinical results with a fusion prototype for mammography and three-dimensional ultrasound with a standard mammography system and a standard ultrasound probe. Acta Radiologica, 2018, 59, 1406-1413.	0.5	10
324	Genetic predictors of chemotherapy-related amenorrhea inÂwomen with breast cancer. Fertility and Sterility, 2019, 112, 731-739.e1.	0.5	10

#	Article	IF	Citations
325	Clinical and analytical validation of Ki-67 in 9069 patients from IBCSG VIIIâ€‰+â€‰IX, BIG1-98 and GeparTrio trial: systematic modulation of interobserver variance in a comprehensive in silico ring trial. Breast Cancer Research and Treatment, 2019, 176, 557-568.	1.1	10
326	Impact of fibroblast growth factor receptor 1 (FGFR1) amplification on the prognosis of breast cancer patients. Breast Cancer Research and Treatment, 2020, 184, 311-324.	1.1	10
327	The association between prenatal alcohol consumption and preschool child stress system disturbance. Developmental Psychobiology, 2021, 63, 687-697.	0.9	10
328	Associations between Genetically Predicted Circulating Protein Concentrations and Endometrial Cancer Risk. Cancers, 2021, 13, 2088.	1.7	10
329	Update Breast Cancer 2021 Part 1 â€“ Prevention and Early Stages. Geburtshilfe Und Frauenheilkunde, 2021, 81, 526-538.	0.8	10
330	Discordance in Human Epidermal Growth Factor Receptor 2 (HER2) Phenotype Between Primary Tumor and Circulating Tumor Cells in Women With HER2-Negative Metastatic Breast Cancer. JCO Precision Oncology, 2017, 1, 1-12.	1.5	9
331	Self-reported Improvement in Side Effects and Quality of Life With Integrative Medicine in Breast Cancer Patients. Integrative Cancer Therapies, 2018, 17, 941-951.	0.8	9
332	Assessment of the additional clinical potential of X-ray dark-field imaging for breast cancer in a preclinical setup. Therapeutic Advances in Medical Oncology, 2020, 12, 175883592095793.	1.4	9
333	A Small Hypoxia Signature Predicted pCR Response to Bevacizumab in the Neoadjuvant GeparQuinto Breast Cancer Trial. Clinical Cancer Research, 2020, 26, 1896-1904.	3.2	9
334	Patterns and Trends of Herbal Medicine Use among Patients with Gynecologic Cancer. Geburtshilfe Und Frauenheilkunde, 2021, 81, 699-707.	0.8	9
335	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. British Journal of Cancer, 2021, 125, 1135-1145.	2.9	9
336	AGO Recommendations for the Diagnosis and Treatment of Patients with Locally Advanced and Metastatic Breast Cancer: Update 2022. Breast Care, 2022, 17, 421-429.	0.8	9
337	Correlates of mammographic density in B-mode ultrasound and real time elastography. European Journal of Cancer Prevention, 2012, 21, 343-349.	0.6	8
338	The UGT1A6_19_GG genotype is a breast cancer risk factor. Frontiers in Genetics, 2013, 4, 104.	1.1	8
339	Update Breast Cancer 2018 (Part 3) â€“ Genomics, Individualized Medicine and Immune Therapies â€“ in the Middle of a New Era: Prevention and Treatment Strategies for Early Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2018, 78, 1110-1118.	0.8	8
340	Differential prognostic relevance of patho-anatomical factors among different tumor-biological subsets of breast cancer: Results from the adjuvant SUCCESS A study. Breast, 2019, 44, 81-89.	0.9	8
341	Breast MRI texture analysis for prediction of BRCA-associated genetic risk. BMC Medical Imaging, 2020, 20, 86.	1.4	8
342	MCM3 is a novel proliferation marker associated with longer survival for patients with tubo-ovarian high-grade serous carcinoma. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2022, 480, 855-871.	1.4	8

#	Article	IF	Citations
343	Breast cancer patientsâ€™ satisfaction with individual therapy goals and treatment in a standardized integrative medicine consultancy service. Archives of Gynecology and Obstetrics, 2018, 298, 147-156.	0.8	7
344	Prenatal Alcohol Exposure and the Facial Phenotype in Adolescents: A Study Based on Meconium Ethyl Glucuronide. Brain Sciences, 2021, 11, 154.	1.1	7
345	Predicting Prognosis of Breast Cancer Patients with Brain Metastases in the BMBC Registryâ€”Comparison of Three Different GPA Prognostic Scores. Cancers, 2021, 13, 844.	1.7	7
346	Association of genomic variants at <scp>PAX8</scp> and <scp>PBX2</scp> with cervical cancer risk. International Journal of Cancer, 2021, 149, 893-900.	2.3	7
347	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
348	A randomized phase II trial to assess the efficacy of paclitaxel and olaparib in comparison to paclitaxel / carboplatin followed by epirubicin / cyclophosphamide as neoadjuvant chemotherapy in patients with HER2-negative early breast cancer and homologous recombination deficiency (HRD): GeparOLA.. Journal of Clinical Oncology, 2016, 34, TPS1096-TPS1096.	0.8	7
349	Analyses of germline variants associated with ovarian cancer survival identify functional candidates at the 1q22 and 19p12 outcome loci. Oncotarget, 2017, 8, 64670-64684.	0.8	7
350	Neurokinin 1 receptor gene polymorphism might be correlated with recurrence rates in endometriosis. Gynecological Endocrinology, 2009, 25, 726-733.	0.7	6
351	Visual pain mapping in endometriosis. Archives of Gynecology and Obstetrics, 2012, 286, 687-693.	0.8	6
352	The postmenopausal hormone replacement therapy-related breast cancer risk is decreased in women carrying the CYP2C19*17 variant. Breast Cancer Research and Treatment, 2012, 131, 347-350.	1.1	6
353	Factors Influencing Decision-Making for or against Adjuvant and Neoadjuvant Chemotherapy in Postmenopausal Hormone Receptor-Positive Breast Cancer Patients in the EvAluate-TM Study. Breast Care, 2016, 11, 315-322.	0.8	6
354	Using automated texture features to determine the probability for masking of a tumor on mammography, but not ultrasound. European Journal of Medical Research, 2017, 22, 30.	0.9	6
355	Preexisting musculoskeletal burden and its development under letrozole treatment in early breast cancer patients. International Journal of Cancer, 2019, 145, 2114-2121.	2.3	6
356	Progression-Free Survival and Overall Survival in Patients with Advanced HER2-Positive Breast Cancer Treated with Trastuzumab Emtansine (T-DM1) after Previous Treatment with Pertuzumab. Cancers, 2020, 12, 3021.	1.7	6
357	Update Breast Cancer 2020 Part 5 â€“ Moving Therapies From Advanced to Early Breast Cancer Patients. Geburtshilfe Und Frauenheilkunde, 2021, 81, 469-480.	0.8	6
358	Update Breast Cancer 2021 Part 2 â€“ Advanced Stages, Long-Term Consequences and Biomarkers. Geburtshilfe Und Frauenheilkunde, 2021, 81, 539-548.	0.8	6
359	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
360	Association of Prenatal Alcohol Exposure and Prenatal Maternal Depression with Offspring Low-Grade Inflammation in Early Adolescence. International Journal of Environmental Research and Public Health, 2021, 18, 7920.	1.2	6

#	Article	IF	Citations
361	Reproducibility of mRNA-Based Testing of ESR1, PGR, ERBB2, and MKI67 Expression in Invasive Breast Cancerâ€"A Europe-Wide External Quality Assessment. Cancers, 2021, 13, 4718.	1.7	6
362	Implementation of an Electronic Patient-Reported Outcome App for Health-Related Quality of Life in Breast Cancer Patients: Evaluation and Acceptability Analysis in a Two-Center Prospective Trial. Journal of Medical Internet Research, 2022, 24, e16128.	2.1	6
363	Rare germline copy number variants (CNVs) and breast cancer risk. Communications Biology, 2022, 5, 65.	2.0	6
364	ABC6 Consensus: Assessment by a Group of German Experts. Breast Care, 2022, 17, 90-100.	0.8	6
365	Update Breast Cancer 2021 Part 5 â€" Advanced Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2022, 82, 215-225.	0.8	6
366	Prevalence of SARS-CoV-2 in Pregnant Women Assessed by RT-PCR in Franconia, Germany: First Results of the SCENARIO Study (SARS-CoV-2 prEvalence in pregNAncy and at biRth In FrancOnia). Geburtshilfe Und Frauenheilkunde, 2022, 82, 226-234.	0.8	6
367	Abstract PD13-06: Neoadjuvant giredestrant (GDC-9545) + palbociclib versus anastrozole + palbociclib in postmenopausal women with estrogen receptor-positive, HER2-negative, untreated early breast cancer: Primary analysis of the randomized, open-label, phase II coopERA breast cancer study. Cancer Research, 2022, 82, PD13-06-PD13-06.	0.4	6
368	7q21-rs6964587 and breast cancer risk: an extended case-control study by the Breast Cancer Association Consortium. Journal of Medical Genetics, 2011, 48, 698-702.	1.5	5
369	Two truncating variants in FANCC and breast cancer risk. Scientific Reports, 2019, 9, 12524.	1.6	5
370	Influence of patient and tumor characteristics on therapy persistence with letrozole in postmenopausal women with advanced breast cancer: results of the prospective observational EvAluate-TM study. BMC Cancer, 2019, 19, 611.	1.1	5
371	Diagnostic Accuracy of Breast Medical Tactile Examiners (MTEs): A Prospective Pilot Study. Breast Care, 2019, 14, 41-47.	0.8	5
372	Risk of postmenopausal hormone therapy and patient history factors for the survival rate in women with endometrial carcinoma. Archives of Gynecology and Obstetrics, 2020, 301, 289-294.	0.8	5
373	RANKL and OPG and their influence on breast volume changes during pregnancy in healthy women. Scientific Reports, 2020, 10, 5171.	1.6	5
374	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
375	Germline BRCA1/2 mutations and severe haematological toxicities in patients with breast cancer treated with neoadjuvant chemotherapy. European Journal of Cancer, 2021, 145, 44-52.	1.3	5
376	Treatment of Patients with Early Breast Cancer: Evidence, Controversies, Consensus. Geburtshilfe Und Frauenheilkunde, 2021, 81, 637-653.	0.8	5
377	Identification of a Locus Near ULK1 Associated With Progression-Free Survival in Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1669-1680.	1.1	5
378	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. Oncotarget, 2016, 7, 69097-69110.	0.8	5

#	Article	IF	Citations
379	Variable Expression of the Disialoganglioside GD2 in Breast Cancer Molecular Subtypes. Cancers, 2021, 13, 5577.	1.7	5
380	Genetic Polymorphisms and Correlation with Treatment-Induced Cardiotoxicity and Prognosis in Patients with Breast Cancer. Clinical Cancer Research, 2022, 28, 1854-1862.	3.2	5
381	MUC1 (CA27.29) before and after Chemotherapy and Prognosis in High-Risk Early Breast Cancer Patients. Cancers, 2022, 14, 1721.	1.7	5
382	Semi-automated De-identification of German Content Sensitive Reports for Big Data Analytics. RoFo Fortschritte Auf Dem Gebiet Der Rontgenstrahlen Und Der Bildgebenden Verfahren, 2017, 189, 661-671.	0.7	4
383	Germline genome-wide association studies in women receiving neoadjuvant chemotherapy with or without bevacizumab. Pharmacogenetics and Genomics, 2018, 28, 147-152.	0.7	4
384	PIWI-Like 1 and PIWI-Like 2 Expression in Breast Cancer. Cancers, 2020, 12, 2742.	1.7	4
385	Treatment Landscape and Prognosis After Treatment with Trastuzumab Emtansine. Geburtshilfe Und Frauenheilkunde, 2020, 80, 1134-1142.	0.8	4
386	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
387	Update Breast Cancer 2021 Part 3 â€“ Current Developments in the Treatment of Early Breast Cancer: Review and Assessment of Specialised Treatment Scenarios by an International Expert Panel. Geburtshilfe Und Frauenheilkunde, 2021, 81, 654-665.	0.8	4
388	Update Breast Cancer 2021 Part 4 â€“ Prevention and Early Stages. Geburtshilfe Und Frauenheilkunde, 2022, 82, 206-214.	0.8	4
389	Abstract PD8-01: Phase 3 SOPHIA study of margetuximab (M) + chemotherapy (CTX) vs trastuzumab (T) + CTX in patients (pts) with HER2+ metastatic breast cancer (MBC) after prior anti-HER2 therapies: Final overall survival (OS) analysis. Cancer Research, 2022, 82, PD8-01-PD8-01.	0.4	4
390	Update Breast Cancer 2018 (Part 4) â€“ Genomics, Individualized Medicine and Immune Therapies â€“ in the Middle of a New Era: Treatment Strategies for Advanced Breast Cancer. Geburtshilfe Und Frauenheilkunde, 2018, 78, 1119-1128.	0.8	3
391	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. International Journal of Molecular Sciences, 2018, 19, 2473.	1.8	3
392	Using Probability for Pathological Complete Response (pCR) as a Decision Support Marker for Neoadjuvant Chemotherapy in HER2 Negative Breast Cancer Patients â€“ a Survey Among Physicians. Geburtshilfe Und Frauenheilkunde, 2018, 78, 707-714.	0.8	3
393	The impact of mammalian target of rapamycin inhibition on bone health in postmenopausal women with hormone receptor-positive advanced breast cancer receiving everolimus plus exemestane in the phase IIIb 4EVER trial. Journal of Bone Oncology, 2019, 14, 100199.	1.0	3
394	Challenges and Opportunities for Real-World Evidence in Metastatic Luminal Breast Cancer. Breast Care, 2021, 16, 108-114.	0.8	3
395	Comparison of C-Reactive Protein in Dried Blood Spots and Saliva of Healthy Adolescents. Frontiers in Immunology, 2021, 12, 795580.	2.2	3
396	Identification of Two Genetic Loci Associated with Leukopenia after Chemotherapy in Patients with Breast Cancer. Clinical Cancer Research, 2022, 28, 3342-3355.	3.2	3

#	Article	IF	Citations
397	Biomarkers in Womenâ€™s Cancers, Gynecology, and Obstetrics. BioMed Research International, 2014, 2014, 1-2.	0.9	2
398	Does anxiety impact the neural processing of child faces in mothers of school-aged children? An ERP study using an emotional Go/NoGo task. Social Neuroscience, 2020, 15, 530-543.	0.7	2
399	Analysis of Oncological Second Opinions in a Certified University Breast and Gynecological Cancer Center Regarding Consensus between the First and Second Opinion and Conformity with the Guidelines. Breast Care, 2021, 16, 291-298.	0.8	2
400	Genetic variations in estrogen and progesterone pathway genes in preeclampsia patients and controls in Bavaria. Archives of Gynecology and Obstetrics, 2021, 303, 897-904.	0.8	2
401	Subgroup of post-neoadjuvant luminal-B tumors assessed by HTG in PENELOPE-B investigating palbociclib in high risk HER2-/HR+ breast cancer with residual disease.. Journal of Clinical Oncology, 2021, 39, 519-519.	0.8	2
402	Comparison of methods for isolation and quantification of circulating cell-free DNA from patients with endometriosis. Reproductive BioMedicine Online, 2021, 43, 788-798.	1.1	2
403	Comprehensive characterization of endometriosis patients and disease patterns in a large clinical cohort. Archives of Gynecology and Obstetrics, 2021, , 1.	0.8	2
404	Breast cancer treatment with everolimus and exemestane for ER+ women: Results of the first interim analysis of the noninterventional trial BRAWO.. Journal of Clinical Oncology, 2014, 32, 578-578.	0.8	2
405	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. Scientific Reports, 2021, 11, 19787.	1.6	2
406	Genetic variants in the genes of the sex steroid hormone metabolism and depressive symptoms during and after pregnancy. Archives of Gynecology and Obstetrics, 2023, 307, 1763-1770.	0.8	2
407	Supportive Infusions in Integrative Breast and Gynecological Oncology â€“ Report on PatientsÊ¼ Satisfaction and Self-reported Effects and Side Effects. Geburtshilfe Und Frauenheilkunde, 2018, 78, 1129-1137.	0.8	1
408	Response to Di Cosimo, Torri, and Porcu. Journal of the National Cancer Institute, 2019, 111, 1234-1235.	3.0	1
409	Heregulin (HRG) assessment for clinical trial eligibility testing in a molecular registry (PRAEGNANT) in Germany. BMC Cancer, 2020, 20, 1091.	1.1	1
410	ABC5 International Consensus Conference on Advanced Breast Cancer, Lisbon, 16 November 2019. Geburtshilfe Und Frauenheilkunde, 2020, 80, 588-600.	0.8	1
411	Genetic variants in the glucocorticoid pathway genes and birth weight. Archives of Gynecology and Obstetrics, 2021, 303, 427-434.	0.8	1
412	Active Participation, Mindâ€“Body Stabilization, and Coping Strategies with Integrative Medicine in Breast Cancer Patients. Integrative Cancer Therapies, 2021, 20, 153473542199010.	0.8	1
413	Abstract PS2-02: Prognostic relevance of the HER2 status of circulating tumor cells in metastatic breast cancer patients screened for participation in the DETECT study program. Cancer Research, 2021, 81, PS2-02-PS2-02.	0.4	1
414	OUP accepted manuscript. Human Molecular Genetics, 2022, , .	1.4	1

#	Article	IF	Citations
415	Occurrence and characteristics of patients with de novo advanced breast cancer according to patient and tumor characteristics â€“ A retrospective analysis of a real world registry. European Journal of Cancer, 2022, 172, 13-21.	1.3	1
416	Response to â€œScreening depression during and after pregnancy using the EPDSâ€: Archives of Gynecology and Obstetrics, 2014, 290, 603-603.	0.8	0
417	Neoadjuvant Treatment of HER2-Positive Breast Cancerâ€”A Review. , 2019, , 95-106.		0
418	Influence of Family History of Breast or Ovarian Cancer on Pathological Complete Response and Long-Term Prognosis in Breast Cancer Patients Treated with Neoadjuvant Chemotherapy. Breast Care, 2021, 16, 254-262.	0.8	0
419	Risikoadaptierte Diagnostik und Therapie. , 2017, , 43-53.		0
420	ZukÃ¼nftige Entwicklungen in der Bildgebung. , 2017, , 201-218.		0
421	A phase II single-arm, multicenter, open-label neoadjuvant study of pembrolizumab in combination with nab-paclitaxel followed by pembrolizumab in combination with epirubicin and cyclophosphamide in patients with triple-negative breast cancer: Neoimmunoboost.. Journal of Clinical Oncology, 2020, 38, e12647-e12647.	0.8	0
422	Update Mammakarzinom 2021 Teil 1 â€“ PrÃvention und frÃ¼he Krankheitsstadien. Senologie - Zeitschrift FÃ¼r Mammadiagnostik Und -therapie, 2021, 18, 377-390.	0.0	0