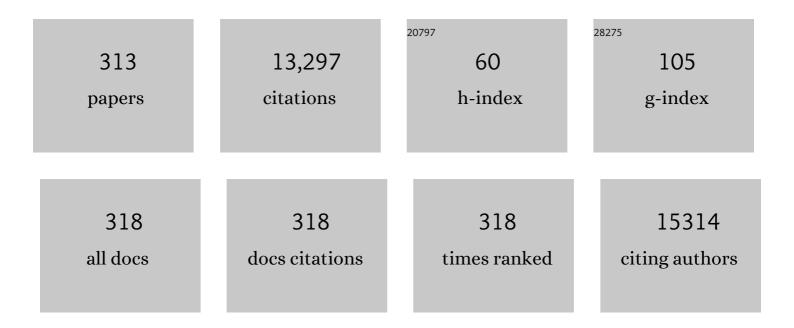
List of Publications by Year in descending order

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ALLISON W KURIAN

#	Article	IF	CITATIONS
1	Influence of payer coverage and outâ€ofâ€pocket costs on ordering of NGS panel tests for hereditary cancer in diverse settings. Journal of Genetic Counseling, 2022, 31, 130-139.	0.9	13
2	Association of Genetic Testing Results With Mortality Among Women With Breast Cancer or Ovarian Cancer. Journal of the National Cancer Institute, 2022, 114, 245-253.	3.0	5
3	Greater financial toxicity relates to greater distress and worse quality of life among breast and gynecologic cancer survivors. Psycho-Oncology, 2022, 31, 9-20.	1.0	29
4	Rare germline copy number variants (CNVs) and breast cancer risk. Communications Biology, 2022, 5, 65.	2.0	6
5	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362.	1.4	23
6	Clinician-Reported Impact of Germline Multigene Panel Testing on Cancer Risk Management Recommendations. JNCI Cancer Spectrum, 2022, 6, .	1.4	1
7	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. Breast Cancer Research, 2022, 24, 2.	2.2	15
8	Breast Cancer Screening Strategies for Women With <i>ATM, CHEK2</i> , and <i>PALB2</i> Pathogenic Variants. JAMA Oncology, 2022, 8, 587.	3.4	36
9	Abstract P2-11-21: Integration of an ancestrally unbiased polygenic risk score with the Tyrer-Cuzick breast cancer risk model. Cancer Research, 2022, 82, P2-11-21-P2-11-21.	0.4	0
10	Polygenic risk scores for prediction of breast cancer risk in Asian populations. Genetics in Medicine, 2022, 24, 586-600.	1.1	27
11	Trends in Annual Surveillance Mammography Participation Among Breast Cancer Survivors From 2004 to 2016. Journal of the National Comprehensive Cancer Network: JNCCN, 2022, 20, 379-386.e9.	2.3	1
12	Weight is More Informative than Body Mass Index for Predicting Postmenopausal Breast Cancer Risk: Prospective Family Study Cohort (ProF-SC). Cancer Prevention Research, 2022, 15, 185-191.	0.7	4
13	Simulation modeling of breast cancer endocrine therapy duration by patient and tumor characteristics. Cancer Medicine, 2022, 11, 297-307.	1.3	2
14	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. Breast Cancer Research, 2022, 24, 27.	2.2	15
15	Relevance of the MHC region for breast cancer susceptibility in Asians. Breast Cancer, 2022, 29, 869-879.	1.3	1
16	Association of illness mindsets with health-related quality of life in cancer survivors Health Psychology, 2022, 41, 389-395.	1.3	4
17	Personalised Risk Prediction in Hereditary Breast and Ovarian Cancer: A Protocol for a Multi-Centre Randomised Controlled Trial. Cancers, 2022, 14, 2716.	1.7	10
18	Breast cancer diagnosis and treatment during the COVID-19 pandemic in a nationwide, insured population. Breast Cancer Research and Treatment, 2022, 194, 475-482.	1.1	14

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19	Harnessing artificial intelligence to automate delineation of volumetric breast cancers from magnetic resonance imaging to improve tumor characterization Journal of Clinical Oncology, 2022, 40, 597-597.	0.8	0
20	Simulation modeling as a tool to support clinical guidelines and care for breast cancer prevention and early detection in high-risk women Journal of Clinical Oncology, 2022, 40, 10525-10525.	0.8	0
21	National claims data analysis of outcomes of hospitalized cancer patients without COVID-19 infection during versus prior to the COVID-19 pandemic Journal of Clinical Oncology, 2022, 40, e18679-e18679.	0.8	0
22	A case-control study of healthcare disparities in sex and gender minority patients with breast cancer Journal of Clinical Oncology, 2022, 40, 6517-6517.	0.8	1
23	Radiomic features quantifying pixel-level characteristics of breast tumors from magnetic resonance imaging predict risk factors in triple-negative breast cancer Journal of Clinical Oncology, 2022, 40, e12612-e12612.	0.8	0
24	A pilot study to increase cascade genetic testing in families with hereditary cancer syndromes Journal of Clinical Oncology, 2022, 40, 10602-10602.	0.8	0
25	Ancestry-specific risk of triple-negative breast cancer (TNBC) associated with germline pathogenic variants (PV) in hereditary cancer (CA) predisposition genes Journal of Clinical Oncology, 2022, 40, 10517-10517.	0.8	0
26	Contributions of screening, early-stage treatment, and metastatic treatment to breast cancer mortality reduction by molecular subtype in U.S. women, 2000-2017 Journal of Clinical Oncology, 2022, 40, 1008-1008.	0.8	3
27	Association of germline genetic testing results with chemotherapy regimens received by women with early-stage breast cancer Journal of Clinical Oncology, 2022, 40, 10518-10518.	0.8	1
28	Constitutional <i>BRCA1</i> methylation and risk of incident triple-negative breast cancer and high-grade serous ovarian cancer Journal of Clinical Oncology, 2022, 40, 10509-10509.	0.8	1
29	Symptoms and survivorship needs differences between "good sleepers―and "bad sleepers―in survivors of breast and gynecologic cancers. Sleep Medicine, 2022, 100, 49-55.	0.8	1
30	Patterns of social media use and associations with psychosocial outcomes among breast and gynecologic cancer survivors. Journal of Cancer Survivorship, 2021, 15, 677-684.	1.5	4
31	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
32	Predicted Chemotherapy Benefit for Breast Cancer Patients With Germline Pathogenic Variants in Cancer Susceptibility Genes. JNCI Cancer Spectrum, 2021, 5, pkaa083.	1.4	3
33	Comparing 5-Year and Lifetime Risks of Breast CancerÂusing the Prospective Family Study Cohort. Journal of the National Cancer Institute, 2021, 113, 785-791.	3.0	13
34	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
35	Impact of the COVID-19 Pandemic on Breast Cancer Mortality in the US: Estimates From Collaborative Simulation Modeling. Journal of the National Cancer Institute, 2021, 113, 1484-1494.	3.0	92
36	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

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37	A Population-Based Study of Genes Previously Implicated in Breast Cancer. New England Journal of Medicine, 2021, 384, 440-451.	13.9	414
38	Association of Risk-Reducing Salpingo-Oophorectomy With Breast Cancer Risk in Women With BRCA1 and BRCA2 Pathogenic Variants. JAMA Oncology, 2021, 7, 585-592.	3.4	16
39	Benchmark Method for Cost Computations Across Health Care Systems: Cost of Care per Patient per Day in Breast Cancer Care. JCO Oncology Practice, 2021, 17, e1403-e1412.	1.4	3
40	Limited English Proficiency and Disparities in Health Care Engagement Among Patients With Breast Cancer. JCO Oncology Practice, 2021, 17, e1837-e1845.	1.4	13
41	Development and Use of Natural Language Processing for Identification of Distant Cancer Recurrence and Sites of Distant Recurrence Using Unstructured Electronic Health Record Data. JCO Clinical Cancer Informatics, 2021, 5, 469-478.	1.0	14
42	Financing of germline testing: implications for availability and access. Molecular Genetics and Metabolism, 2021, 132, S330-S331.	0.5	0
43	Treatment and Monitoring Variability in US Metastatic Breast Cancer Care. JCO Clinical Cancer Informatics, 2021, 5, 600-614.	1.0	5
44	Time Trends in Receipt of Germline Genetic Testing and Results for Women Diagnosed With Breast Cancer or Ovarian Cancer, 2012-2019. Journal of Clinical Oncology, 2021, 39, 1631-1640.	0.8	62
45	Multicancer hereditary syndrome testing: Genetic counselors' perspectives Journal of Clinical Oncology, 2021, 39, 10594-10594.	0.8	0
46	Cancer-specific mortality associated with germline genetic testing results among women with breast cancer or ovarian cancer treated with chemotherapy Journal of Clinical Oncology, 2021, 39, 10517-10517.	0.8	0
47	Twenty-one-gene recurrence score (RS) in germline (g)CHEK2 mutation-associated versus sporadic breast cancers (BC): A multi-site case-control study Journal of Clinical Oncology, 2021, 39, 10531-10531.	0.8	0
48	Impact of disruptions in breast cancer control due to the COVID-19 pandemic on breast cancer mortality in the United States: Estimates from collaborative simulation modeling Journal of Clinical Oncology, 2021, 39, 6562-6562.	0.8	0
49	Breast cancer screening for carriers of ATM, CHEK2, and PALB2 pathogenic variants: A comparative modeling analysis Journal of Clinical Oncology, 2021, 39, 10500-10500.	0.8	0
50	Weakly supervised temporal model for prediction of breast cancer distant recurrence. Scientific Reports, 2021, 11, 9461.	1.6	11
51	A simulation model-based clinical decision tool to guide personalized treatment based on individual characteristics: Does 21-gene recurrence score assay testing change decisions?. Journal of Clinical Oncology, 2021, 39, e12507-e12507.	0.8	0
52	Comprehensive Breast Cancer Risk Assessment for <i>CHEK2</i> and <i>ATM</i> Pathogenic Variant Carriers Incorporating a Polygenic Risk Score and the Tyrer-Cuzick Model. JCO Precision Oncology, 2021, 5, 1073-1081.	1.5	9
53	Receipt of guidelineâ€concordant care among young adult women with breast cancer. Cancer, 2021, 127, 3325-3333.	2.0	3
54	Tobacco Smoking and Risk of Second Primary Lung Cancer. Journal of Thoracic Oncology, 2021, 16, 968-979.	0.5	54

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55	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
56	Performance of the IBIS/Tyrerâ€Cuzick model of breast cancer risk by race and ethnicity in the Women's Health Initiative. Cancer, 2021, 127, 3742-3750.	2.0	21
57	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
58	Widening cancer care disparities in the adoption of telemedicine during COVID 19: who is left behind?. Gynecologic Oncology, 2021, 162, S23.	0.6	2
59	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. British Journal of Cancer, 2021, 125, 1135-1145.	2.9	9
60	Genetic insights into biological mechanisms governing human ovarian ageing. Nature, 2021, 596, 393-397.	13.7	183
61	Development of a Mobile Health App (TOGETHERCare) to Reduce Cancer Care Partner Burden: Product Design Study. JMIR Formative Research, 2021, 5, e22608.	0.7	9
62	Development and Validation of a Simulation Model–Based Clinical Decision Tool: Identifying Patients Where 21-Gene Recurrence Score Testing May Change Decisions. Journal of Clinical Oncology, 2021, 39, 2893-2902.	0.8	7
63	The Impact of COVID-19 on Patients With Cancer. American Journal of Clinical Oncology: Cancer Clinical Trials, 2021, 44, 580-587.	0.6	26
64	Impact of Low-Dose Computed Tomography Screening for Primary Lung Cancer on Subsequent Risk of Brain Metastasis. Journal of Thoracic Oncology, 2021, 16, 1479-1489.	0.5	2
65	Multicancer hereditary syndrome testing: Genetic counselors' perspectives Journal of Clinical Oncology, 2021, 39, 106-106.	0.8	1
66	Germline Pathogenic Variants in the Ataxia Telangiectasia Mutated (<i>ATM</i>) Gene are Associated with High and Moderate Risks for Multiple Cancers. Cancer Prevention Research, 2021, 14, 433-440.	0.7	68
67	Integrating Clinical and Polygenic Factors to Predict Breast Cancer Risk in Women Undergoing Genetic Testing. JCO Precision Oncology, 2021, 5, 307-316.	1.5	18
68	Genetic/Familial High-Risk Assessment: Breast, Ovarian, and Pancreatic, Version 2.2021, NCCN Clinical Practice Guidelines in Oncology. Journal of the National Comprehensive Cancer Network: JNCCN, 2021, 19, 77-102.	2.3	498
69	Prevalence of Lynch syndrome in women with mismatch repairâ€deficient ovarian cancer. Cancer Medicine, 2021, 10, 1012-1017.	1.3	12
70	Psychosocial outcomes following germline multigene panel testing in an ethnically and economically diverse cohort of patients. Cancer, 2021, 127, 1275-1285.	2.0	21
71	Reply to Ritzwoller et al. JCO Clinical Cancer Informatics, 2021, 5, 1026-1027.	1.0	0
72	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. Scientific Reports, 2021, 11, 19787.	1.6	2

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73	Multiple imputation with missing data indicators. Statistical Methods in Medical Research, 2021, 30, 2685-2700.	0.7	30
74	Germline Pathogenic Variants in Cancer Predisposition Genes Among Women With Invasive Lobular Carcinoma of the Breast. Journal of Clinical Oncology, 2021, 39, 3918-3926.	0.8	22
75	Racial/Ethnic Disparities in Survival after Breast Cancer Diagnosis by Estrogen and Progesterone Receptor Status: A Pooled Analysis. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 351-363.	1.1	7
76	Racial/ethnic differences in cancer diagnosed after metastasis: absolute burden and deaths potentially avoidable through earlier detection. Cancer Epidemiology Biomarkers and Prevention, 2021, , cebp.0823.2021.	1.1	7
77	Association of Family Cancer History With Pathogenic Variants in Specific Breast Cancer Susceptibility Genes. JCO Precision Oncology, 2021, 5, 1853-1859.	1.5	2
78	Recreational Physical Activity and Outcomes After Breast Cancer in Women at High Familial Risk. JNCI Cancer Spectrum, 2021, 5, pkab090.	1.4	1
79	Decision Making About Genetic Testing Among Women With a Personal and Family History of Breast Cancer. JCO Oncology Practice, 2020, 16, e37-e55.	1.4	16
80	Magnitude of reduction in risk of second contralateral breast cancer with bilateral mastectomy in patients with breast cancer: Data from California, 1998 through 2015. Cancer, 2020, 126, 958-970.	2.0	11
81	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. Nature Communications, 2020, 11, 3833.	5.8	88
82	Association of a Polygenic Risk Score With Breast Cancer Among Women Carriers of High- and Moderate-Risk Breast Cancer Genes. JAMA Network Open, 2020, 3, e208501.	2.8	79
83	Pathogenic Variants in Breast Cancer Susceptibility Genes in Older Women—Reply. JAMA - Journal of the American Medical Association, 2020, 324, 397.	3.8	Ο
84	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. American Journal of Human Genetics, 2020, 107, 837-848.	2.6	39
85	Yield and Utility of Germline Testing Following Tumor Sequencing in Patients With Cancer. JAMA Network Open, 2020, 3, e2019452.	2.8	76
86	Projected Reductions in Absolute Cancer–Related Deaths from Diagnosing Cancers Before Metastasis, 2006–2015. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 895-902.	1.1	36
87	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. Scientific Reports, 2020, 10, 9688.	1.6	2
88	Development and Validation of a Clinical Polygenic Risk Score to Predict Breast Cancer Risk. JCO Precision Oncology, 2020, 4, 585-592.	1.5	41
89	Hospital Characteristics and Breast Cancer Survival in the California Breast Cancer Survivorship Consortium. JCO Oncology Practice, 2020, 16, e517-e528.	1.4	6
90	Prevalence of Pathogenic Variants in Cancer Susceptibility Genes Among Women With Postmenopausal Breast Cancer. JAMA - Journal of the American Medical Association, 2020, 323, 995.	3.8	26

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91	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. Nature Communications, 2020, 11, 1217.	5.8	46
92	Characterization of the Cancer Spectrum in Men With Germline <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. JAMA Oncology, 2020, 6, 1218.	3.4	48
93	Emerging Opportunity of Cascade Genetic Testing for Population-Wide Cancer Prevention and Control. Journal of Clinical Oncology, 2020, 38, 1371-1374.	0.8	18
94	Health Disparities in Germline Genetic Testing for Cancer Susceptibility. Current Breast Cancer Reports, 2020, 12, 51-58.	0.5	6
95	Association of Germline Genetic Testing Results With Locoregional and Systemic Therapy in Patients With Breast Cancer. JAMA Oncology, 2020, 6, e196400.	3.4	32
96	Insights From a Temporal Assessment of Increases in US Private Payer Coverage of Tumor Sequencing From 2015 to 2019. Value in Health, 2020, 23, 551-558.	0.1	9
97	A case of a transâ€masculine patient receiving testosterone with a history of estrogen receptorâ€positive breast cancer. Breast Journal, 2020, 26, 1888-1889.	0.4	5
98	Abstract P3-07-01: Breast cancer-specific mortality (BCSM) in patients age 50 years or younger with node-positive (N+) breast cancer (BC) treated based on the 21-gene assay in clinical practice. Cancer Research, 2020, 80, P3-07-01-P3-07-01.	0.4	2
99	Performance of the IBIS/Tyrer-Cuzick (TC) Model by race/ethnicity in the Women's Health Initiative Journal of Clinical Oncology, 2020, 38, 1503-1503.	0.8	2
100	NCCN Guidelines Insights: Genetic/Familial High-Risk Assessment: Breast, Ovarian, and Pancreatic, Version 1.2020. Journal of the National Comprehensive Cancer Network: JNCCN, 2020, 18, 380-391.	2.3	314
101	Abstract P6-08-02: 21-gene recurrence score results according to germline pathogenic variants inBRCA1, BRCA2, PALB2, ATM, CHEK2and Lynch Syndrome genes. , 2020, , .		0
102	Linking insurance claims across time to characterize treatment, monitoring, and end-of-life care in metastatic breast cancer Journal of Clinical Oncology, 2020, 38, 7063-7063.	0.8	0
103	Comprehensive breast cancer (BC) risk assessment for CHEK2 carriers incorporating a polygenic risk score (PRS) and the Tyrer-Cuzick (TC) model Journal of Clinical Oncology, 2020, 38, 1504-1504.	0.8	0
104	Clinicopathologic features of invasive breast cancer (BC) diagnosed in carriers of germline <i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> pathogenic variants Journal of Clinical Oncology, 2020, 38, 1549-1549.	0.8	2
105	Real-world outcomes of patients with metastatic breast cancer (BC) treated with osteoclast inhibitors (Ols) Journal of Clinical Oncology, 2020, 38, e19314-e19314.	0.8	0
106	Development and validation of natural language processing (NLP) algorithm for detection of distant versus local breast cancer recurrence and metastatic site Journal of Clinical Oncology, 2020, 38, 2043-2043.	0.8	1
107	Abstract IA50: Genetic testing, treatment and mortality after diagnosis of breast cancer or ovarian cancer: The SEER-GeneLINK Initiative. , 2020, , .		0
108	Abstract 2033: Reducing cancer caregiver burden: A user-centered design approach for an mHealth app.		0

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109	Reply to Residual confounding threatens the validity of observational studies on breast cancer local therapy. Cancer, 2020, 126, 2317-2318.	2.0	Ο
110	Abstract P5-03-02: Cancer risks associated with pathogenic variants in the ataxia telangiectasia mutated (ATM) gene. , 2020, , .		0
111	Abstract P6-08-07: Polygenic breast cancer risk modification in carriers of high and intermediate risk gene mutations. , 2020, , .		Ο
112	Trends in germline genetic testing and results into survivorship for women diagnosed with breast cancer or ovarian cancer, 2013 to 2017 Journal of Clinical Oncology, 2020, 38, 273-273.	0.8	0
113	Genomic landscape of ductal carcinoma in situ and association with progression. Breast Cancer Research and Treatment, 2019, 178, 307-316.	1.1	17
114	Chromatin Remodeling in Response to BRCA2-Crisis. Cell Reports, 2019, 28, 2182-2193.e6.	2.9	6
115	Multicenter Prospective Cohort Study of the Diagnostic Yield and Patient Experience of Multiplex Gene Panel Testing For Hereditary Cancer Risk. JCO Precision Oncology, 2019, 3, 1-12.	1.5	23
116	Patient-clinician interactions and disparities in breast cancer care: the equality in breast cancer care study. Journal of Cancer Survivorship, 2019, 13, 968-980.	1.5	14
117	Using natural language processing to construct a metastatic breast cancer cohort from linked cancer registry and electronic medical records data. JAMIA Open, 2019, 2, 528-537.	1.0	40
118	Re-evaluating genetic variants identified in candidate gene studies of breast cancer risk using data from nearly 280,000 women of Asian and European ancestry. EBioMedicine, 2019, 48, 203-211.	2.7	14
119	Simulation Modeling to Extend Clinical Trials of Adjuvant Chemotherapy Guided by a 21-Gene Expression Assay in Early Breast Cancer. JNCI Cancer Spectrum, 2019, 3, pkz062.	1.4	2
120	Primary care provider–reported involvement in breast cancer treatment decisions. Cancer, 2019, 125, 1815-1822.	2.0	12
121	Is Breast Cancer in Asian and Asian American Women a Different Disease?. Journal of the National Cancer Institute, 2019, 111, 1243-1244.	3.0	17
122	Response to Peshkin, Isaacs, and Schwartz. Journal of the National Cancer Institute, 2019, 111, 874-874.	3.0	0
123	Genetic Testing and Results in a Population-Based Cohort of Breast Cancer Patients and Ovarian Cancer Patients. Journal of Clinical Oncology, 2019, 37, 1305-1315.	0.8	266
124	Automatic inference of BI-RADS final assessment categories from narrative mammography report findings. Journal of Biomedical Informatics, 2019, 92, 103137.	2.5	14
125	Comparative effectiveness of first-line nab-paclitaxel versus paclitaxel monotherapy in triple-negative breast cancer. Journal of Comparative Effectiveness Research, 2019, 8, 1173-1185.	0.6	3
126	Can precision medicine help achieve the goal of reducing care when the risks exceed the benefits?. Personalized Medicine, 2019, 16, 365-367.	0.8	0

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127	Natural Language Processing Approaches to Detect the Timeline of Metastatic Recurrence of Breast Cancer. JCO Clinical Cancer Informatics, 2019, 3, 1-12.	1.0	43
128	Guidelines Do Not Proscribe Surgeons Performing Genetic Testing—Reply. JAMA Surgery, 2019, 154, 269.	2.2	0
129	Distribution of global health measures from routinely collected PROMIS surveys in patients with breast cancer or prostate cancer. Cancer, 2019, 125, 943-951.	2.0	15
130	Cascade Genetic Testing of Relatives for Hereditary Cancer Risk: Results of an Online Initiative. Journal of the National Cancer Institute, 2019, 111, 95-98.	3.0	81
131	Preventive surgery after multiplex genetic panel testing (MGPT) Journal of Clinical Oncology, 2019, 37, 1525-1525.	0.8	1
132	Breast cancer treatment according to pathogenic variants in cancer susceptibility genes in a population-based cohort Journal of Clinical Oncology, 2019, 37, 560-560.	0.8	2
133	Oncotype DX DCIS use and clinical utility: A SEER population-based study Journal of Clinical Oncology, 2019, 37, e12046-e12046.	0.8	1
134	Uptake of the 21-Gene Assay Among Women With Node-Positive, Hormone Receptorâ^'Positive Breast Cancer. Journal of the National Comprehensive Cancer Network: JNCCN, 2019, 17, 662-668.	2.3	14
135	Prevalence and penetrance of breast cancer-associated mutations identified by multiple-gene sequencing in the Women's Health Initiative Journal of Clinical Oncology, 2019, 37, 1513-1513.	0.8	0
136	Radiomics features to identify distinct subtypes of triple-negative breast cancers Journal of Clinical Oncology, 2019, 37, 3069-3069.	0.8	0
137	Use, attitudes, and perceptions of tumor genomic testing: Survey of TAPUR physicians Journal of Clinical Oncology, 2019, 37, 6531-6531.	0.8	1
138	Differences among Asian/Asian American, and Caucasian breast and gynecologic cancer patient-reported survivorship needs, symptoms, and illness mindsets (N=220) Journal of Clinical Oncology, 2019, 37, 11620-11620.	0.8	0
139	Adherence to breast cancer treatment guidelines according to pathogenic variants in cancer susceptibility genes in a population-based cohort Journal of Clinical Oncology, 2019, 37, 34-34.	0.8	0
140	Patient Experiences and Clinician Views on the Role of Radiation Therapy for Ductal Carcinoma In Situ. International Journal of Radiation Oncology Biology Physics, 2018, 100, 1237-1245.	0.4	10
141	Association of Screening and Treatment With Breast Cancer Mortality by Molecular Subtype in US Women, 2000-2012. JAMA - Journal of the American Medical Association, 2018, 319, 154.	3.8	209
142	Recent Trends in Chemotherapy Use and Oncologists' Treatment Recommendations for Early-Stage Breast Cancer. Journal of the National Cancer Institute, 2018, 110, 493-500.	3.0	50
143	Intratumoral Spatial Heterogeneity at Perfusion MR Imaging Predicts Recurrence-free Survival in Locally Advanced Breast Cancer Treated with Neoadjuvant Chemotherapy. Radiology, 2018, 288, 26-35.	3.6	102
144	Common Model Inputs Used in CISNET Collaborative Breast Cancer Modeling. Medical Decision Making, 2018, 38, 9S-23S.	1.2	37

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145	Differences in Breast Cancer Survival by Molecular Subtypes in the United States. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 619-626.	1.1	341
146	Higher Absolute Lymphocyte Counts Predict Lower Mortality from Early-Stage Triple-Negative Breast Cancer. Clinical Cancer Research, 2018, 24, 2851-2858.	3.2	65
147	What Factors Influence Women's Perceptions of their Systemic Recurrence Risk after Breast Cancer Treatment?. Medical Decision Making, 2018, 38, 95-106.	1.2	8
148	Racial/ethnic differences in multiple-gene sequencing results for hereditary cancer risk. Genetics in Medicine, 2018, 20, 234-239.	1.1	131
149	Association of Germline Genetic Test Type and Results With Patient Cancer Worry After Diagnosis of Breast Cancer. JCO Precision Oncology, 2018, 2018, 1-8.	1.5	9
150	Gaps in Receipt of Clinically Indicated Genetic Counseling After Diagnosis of Breast Cancer. Journal of Clinical Oncology, 2018, 36, 1218-1224.	0.8	59
151	Rapid detection of <i>BRCA1/2</i> recurrent mutations in Chinese breast and ovarian cancer patients with multiplex SNaPshot genotyping panels. Oncotarget, 2018, 9, 7832-7843.	0.8	9
152	Pathogenic Variants in Less Familiar Cancer Susceptibility Genes: What Happens After Genetic Testing?. JCO Precision Oncology, 2018, 2, 1-10.	1.5	7
153	Change in Survival in Metastatic Breast Cancer with Treatment Advances: Meta-Analysis and Systematic Review. JNCI Cancer Spectrum, 2018, 2, pky062.	1.4	199
154	Cancer Risk Estimates for Study of Multiple-Gene Testing After Diagnosis of Breast Cancer—Reply. JAMA Oncology, 2018, 4, 1788.	3.4	1
155	Patient communication of cancer genetic test results in a diverse population. Translational Behavioral Medicine, 2018, 8, 85-94.	1.2	34
156	Knowledge regarding and patterns of genetic testing in patients newly diagnosed with breast cancer participating in the iCanDecide trial. Cancer, 2018, 124, 4000-4009.	2.0	6
157	Can We Use Survival Data from Cancer Registries to Learn about Disease Recurrence? The Case of Breast Cancer. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 1332-1341.	1.1	28
158	Association of Attending Surgeon With Variation in the Receipt of Genetic Testing After Diagnosis of Breast Cancer. JAMA Surgery, 2018, 153, 909.	2.2	22
159	Unmet need for clinician engagement regarding financial toxicity after diagnosis of breast cancer. Cancer, 2018, 124, 3668-3676.	2.0	118
160	From the Past to the Present: Insurer Coverage Frameworks for Next-Generation Tumor Sequencing. Value in Health, 2018, 21, 1062-1068.	0.1	19
161	Rising rates of bilateral mastectomy with reconstruction following neoadjuvant chemotherapy. International Journal of Cancer, 2018, 143, 3262-3272.	2.3	18
162	Measuring serum melatonin in postmenopausal women: Implications for epidemiologic studies and breast cancer studies. PLoS ONE, 2018, 13, e0195666.	1.1	5

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163	Uptake, Results, and Outcomes of Germline Multiple-Gene Sequencing After Diagnosis of Breast Cancer. JAMA Oncology, 2018, 4, 1066.	3.4	146
164	Macrophages Promote Circulating Tumor Cell–Mediated Local Recurrence following Radiotherapy in Immunosuppressed Patients. Cancer Research, 2018, 78, 4241-4252.	0.4	36
165	Unmet need for clinician engagement about financial toxicity after diagnosis of breast cancer Journal of Clinical Oncology, 2018, 36, 10080-10080.	0.8	1
166	Computing the cost of care per day of breast cancer survivor care Journal of Clinical Oncology, 2018, 36, 10-10.	0.8	1
167	Promoting colorectal cancer (CRC) screening after multiplex genetic testing and genetic counseling Journal of Clinical Oncology, 2018, 36, 1582-1582.	0.8	0
168	Genetic testing and results in population-based breast cancer patients and ovarian cancer patients Journal of Clinical Oncology, 2018, 36, 1578-1578.	0.8	0
169	Promoting breast cancer screening after multiplex genetic panel testing (MGPT) and genetic counseling Journal of Clinical Oncology, 2018, 36, 1581-1581.	0.8	0
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