# Olufunmilayo I Olopade

## List of Publications by Citations

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12,293 107 223 53 h-index g-index citations papers 15,381 5.82 254 9.3 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
223	Comprehensive Molecular Portraits of Invasive Lobular Breast Cancer. <i>Cell</i> , <b>2015</b> , 163, 506-19	56.2	1055
222	Association of risk-reducing surgery in BRCA1 or BRCA2 mutation carriers with cancer risk and mortality. <i>JAMA - Journal of the American Medical Association</i> , <b>2010</b> , 304, 967-75	27.4	993
221	Bilateral prophylactic mastectomy reduces breast cancer risk in BRCA1 and BRCA2 mutation carriers: the PROSE Study Group. <i>Journal of Clinical Oncology</i> , <b>2004</b> , 22, 1055-62	2.2	924
220	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94	50.4	643
219	Wnt/beta-catenin pathway activation is enriched in basal-like breast cancers and predicts poor outcome. <i>American Journal of Pathology</i> , <b>2010</b> , 176, 2911-20	5.8	361
218	Breast cancer risk associated with BRCA1 and BRCA2 in diverse populations. <i>Nature Reviews Cancer</i> , <b>2007</b> , 7, 937-48	31.3	339
217	Association of type and location of BRCA1 and BRCA2 mutations with risk of breast and ovarian cancer. <i>JAMA - Journal of the American Medical Association</i> , <b>2015</b> , 313, 1347-61	27.4	286
216	Population differences in breast cancer: survey in indigenous African women reveals over-representation of triple-negative breast cancer. <i>Journal of Clinical Oncology</i> , <b>2009</b> , 27, 4515-21	2.2	284
215	Common breast cancer-predisposition alleles are associated with breast cancer risk in BRCA1 and BRCA2 mutation carriers. <i>American Journal of Human Genetics</i> , <b>2008</b> , 82, 937-48	11	218
214	Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003212	6	209
213	RAD51 135G>C modifies breast cancer risk among BRCA2 mutation carriers: results from a combined analysis of 19 studies. <i>American Journal of Human Genetics</i> , <b>2007</b> , 81, 1186-200	11	204
212	Proliferating macrophages associated with high grade, hormone receptor negative breast cancer and poor clinical outcome. <i>Breast Cancer Research and Treatment</i> , <b>2011</b> , 128, 703-711	4.4	193
211	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer.  Nature Genetics, 2017, 49, 680-691	36.3	190
<b>21</b> 0	Genetic testing in an ethnically diverse cohort of high-risk women: a comparative analysis of BRCA1 and BRCA2 mutations in American families of European and African ancestry. <i>JAMA - Journal of the American Medical Association</i> , <b>2005</b> , 294, 1925-33	27.4	189
209	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer.  Nature Genetics, 2017, 49, 1767-1778	36.3	186
208	A perfect storm: How tumor biology, genomics, and health care delivery patterns collide to create a racial survival disparity in breast cancer and proposed interventions for change. <i>Ca-A Cancer Journal for Clinicians</i> , <b>2015</b> , 65, 221-38	220.7	181
207	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 164-71	36.3	177

#### (2018-2008)

206	Targeting health disparities: a model linking upstream determinants to downstream interventions. <i>Health Affairs</i> , <b>2008</b> , 27, 339-49	7	177
205	Assembly of a pan-genome from deep sequencing of 910 humans of African descent. <i>Nature Genetics</i> , <b>2019</b> , 51, 30-35	36.3	153
204	Comparison of Breast Cancer Molecular Features and Survival by African and European Ancestry in The Cancer Genome Atlas. <i>JAMA Oncology</i> , <b>2017</b> , 3, 1654-1662	13.4	146
203	Mutational spectrum in a worldwide study of 29,700 families with BRCA1 or BRCA2 mutations. <i>Human Mutation</i> , <b>2018</b> , 39, 593-620	4.7	138
202	Advances in breast cancer: pathways to personalized medicine. Clinical Cancer Research, 2008, 14, 7988	<b>-92</b> .9	134
201	Cancer Risks Associated With Germline Pathogenic Variants: An International Study of 524 Families. Journal of Clinical Oncology, <b>2020</b> , 38, 674-685	2.2	133
200	Breast cancer genetics in African Americans. <i>Cancer</i> , <b>2003</b> , 97, 236-45	6.4	131
199	Disparities in genetic testing: thinking outside the BRCA box. Journal of Clinical Oncology, 2006, 24, 219	07 <u>&gt;2</u> 03	126
198	ECatenin is required for the tumorigenic behavior of triple-negative breast cancer cells. <i>PLoS ONE</i> , <b>2015</b> , 10, e0117097	3.7	111
197	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , <b>2018</b> , 50, 968-978	36.3	101
196	Clinical, morphologic, and cytogenetic characteristics of patients with lymphoid malignancies characterized by both t(14;18)(q32;q21) and t(8;14)(q24;q32) or t(8;22)(q24;q11). <i>Genes Chromosomes and Cancer</i> , <b>1990</b> , 2, 147-58	5	97
195	Inherited predisposition to breast cancer among African American women. <i>Breast Cancer Research and Treatment</i> , <b>2015</b> , 149, 31-9	4.4	95
194	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , <b>2016</b> , 48, 374-86	36.3	93
193	A continuum of admixture in the Western Hemisphere revealed by the African Diaspora genome. <i>Nature Communications</i> , <b>2016</b> , 7, 12522	17.4	90
192	Inherited mutations in cancer susceptibility genes are common among survivors of breast cancer who develop therapy-related leukemia. <i>Cancer</i> , <b>2016</b> , 122, 304-11	6.4	89
191	Prevalence of BRCA1 and BRCA2 mutations among clinic-based African American families with breast cancer. <i>Human Genetics</i> , <b>2000</b> , 107, 186-91	6.3	89
190	Effect of Oophorectomy on Survival After Breast Cancer in BRCA1 and BRCA2 Mutation Carriers. JAMA Oncology, <b>2015</b> , 1, 306-13	13.4	81
189	HIF-2[promotes conversion to a stem cell phenotype and induces chemoresistance in breast cancer cells by activating Wnt and Notch pathways. <i>Journal of Experimental and Clinical Cancer Research</i> 2018, 37, 256	12.8	77

188	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , <b>2020</b> , 52, 572-581	36.3	76
187	The Breast-Thyroid Cancer Link: A Systematic Review and Meta-analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 231-8	4	68
186	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , <b>2016</b> , 108,	9.7	65
185	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , <b>2016</b> , 7, 11375	17.4	64
184	FANCF methylation contributes to chemoselectivity in ovarian cancer. Cancer Cell, 2003, 3, 417-20	24.3	64
183	A genome-wide association study of breast cancer in women of African ancestry. <i>Human Genetics</i> , <b>2013</b> , 132, 39-48	6.3	63
182	High prevalence of BRCA1 and BRCA2 mutations in unselected Nigerian breast cancer patients. <i>International Journal of Cancer</i> , <b>2012</b> , 131, 1114-23	7.5	63
181	Genetic and functional analysis of CHEK2 (CHK2) variants in multiethnic cohorts. <i>International Journal of Cancer</i> , <b>2007</b> , 121, 2661-7	7.5	62
180	Racial Differences in PAM50 Subtypes in the Carolina Breast Cancer Study. <i>Journal of the National Cancer Institute</i> , <b>2018</b> , 110,	9.7	62
179	Building capacity for sustainable research programmes for cancer in Africa. <i>Nature Reviews Clinical Oncology</i> , <b>2014</b> , 11, 251-9	19.4	61
178	Development of a tiered and binned genetic counseling model for informed consent in the era of multiplex testing for cancer susceptibility. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 485-92	8.1	60
177	Computerized analysis of mammographic parenchymal patterns for breast cancer risk assessment: feature selection. <i>Medical Physics</i> , <b>2000</b> , 27, 4-12	4.4	60
176	Molecular-cytogenetic analysis of HER-2/neu gene in BRCA1-associated breast cancers. <i>Cancer Research</i> , <b>2002</b> , 62, 1481-8	10.1	60
175	Evaluation of 19 susceptibility loci of breast cancer in women of African ancestry. <i>Carcinogenesis</i> , <b>2012</b> , 33, 835-40	4.6	59
174	Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 15	8.3	58
173	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , <b>2020</b> , 52, 56-73	36.3	56
172	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , <b>2016</b> , 7, 12675	17.4	53
171	Efficacy of risk-reducing salpingo-oophorectomy in women with BRCA-1 and BRCA-2 mutations. <i>Breast Journal</i> , <b>2004</b> , 10 Suppl 1, S5-9	1.2	53

170	Identification of a circulating microRNA signature to distinguish recurrence in breast cancer patients. <i>Oncotarget</i> , <b>2016</b> , 7, 55231-55248	3.3	53	
169	Inherited Breast Cancer in Nigerian Women. <i>Journal of Clinical Oncology</i> , <b>2018</b> , 36, 2820-2825	2.2	51	
168	Codeletion of CDKN2 and MTAP genes in a subset of non-Hodgkinß lymphoma may be associated with histologic transformation from low-grade to diffuse large-cell lymphoma. <i>Genes Chromosomes and Cancer</i> , <b>1998</b> , 22, 72-78	5	48	
167	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , <b>2019</b> , 10, 1741	17.4	47	
166	Associations of common breast cancer susceptibility alleles with risk of breast cancer subtypes in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 3416	8.3	46	
165	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , <b>2019</b> , 10, 431	17.4	45	
164	DNA repair deficiency biomarkers and the 70-gene ultra-high risk signature as predictors of veliparib/carboplatin response in the I-SPY 2 breast cancer trial. <i>Npj Breast Cancer</i> , <b>2017</b> , 3, 31	7.8	45	
163	Characterization of Nigerian breast cancer reveals prevalent homologous recombination deficiency and aggressive molecular features. <i>Nature Communications</i> , <b>2018</b> , 9, 4181	17.4	45	
162	Metabolically activated adipose tissue macrophages link obesity to triple-negative breast cancer. Journal of Experimental Medicine, <b>2019</b> , 216, 1345-1358	16.6	42	
161	Association of Event-Free and Distant Recurrence-Free Survival With Individual-Level Pathologic Complete Response in Neoadjuvant Treatment of Stages 2 and 3 Breast Cancer: Three-Year Follow-up Analysis for the I-SPY2 Adaptively Randomized Clinical Trial. <i>JAMA Oncology</i> , <b>2020</b> , 6, 1355-1.	13.4 <b>362</b>	42	
160	Prediction of BRCA Mutations Using the BRCAPRO Model in Clinic-Based African American, Hispanic, and Other Minority Families in the United States. <i>Journal of Clinical Oncology</i> , <b>2009</b> , 27, 1184-9	9 <b>ð</b> .2	40	
159	Obesity and height in urban Nigerian women with breast cancer. <i>Annals of Epidemiology</i> , <b>2003</b> , 13, 455-	6 <b>6</b> .4	40	
158	Therapeutic plasma exchange for the acute management of the catastrophic antiphospholipid syndrome: beta(2)-glycoprotein I antibodies as a marker of response to therapy. <i>Journal of Clinical Apheresis</i> , <b>1999</b> , 14, 171-6	3.2	38	
157	Mutation spectrum and risk of colorectal cancer in African American families with Lynch syndrome. <i>Gastroenterology</i> , <b>2015</b> , 149, 1446-53	13.3	37	
156	Exatenin regulates c-Myc and CDKN1A expression in breast cancer cells. <i>Molecular Carcinogenesis</i> , <b>2016</b> , 55, 431-9	5	37	
155	Naturally occurring BRCA2 alternative mRNA splicing events in clinically relevant samples. <i>Journal of Medical Genetics</i> , <b>2016</b> , 53, 548-58	5.8	37	
154	Association study in African-admixed populations across the Americas recapitulates asthma risk loci in non-African populations. <i>Nature Communications</i> , <b>2019</b> , 10, 880	17.4	36	
153	Recurrent BRCA1 and BRCA2 mutations in breast cancer patients of African ancestry. <i>Breast Cancer Research and Treatment</i> , <b>2012</b> , 134, 889-94	4.4	35	

152	Effect of stove intervention on household air pollution and the respiratory health of women and children in rural Nigeria. <i>Air Quality, Atmosphere and Health</i> , <b>2013</b> , 6, 553-561	5.6	34
151	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , <b>2020</b> , 22, 1653-1666	8.1	34
150	Genome-wide association studies in women of African ancestry identified 3q26.21 as a novel susceptibility locus for oestrogen receptor negative breast cancer. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 4835-4846	5.6	34
149	DNA glycosylases involved in base excision repair may be associated with cancer risk in BRCA1 and BRCA2 mutation carriers. <i>PLoS Genetics</i> , <b>2014</b> , 10, e1004256	6	33
148	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , <b>2018</b> , 78, 5419-5430	10.1	32
147	Radiogenomics of breast cancer using dynamic contrast enhanced MRI and gene expression profiling. <i>Cancer Imaging</i> , <b>2019</b> , 19, 48	5.6	32
146	Relationships between computer-extracted mammographic texture pattern features and BRCA1/2 mutation status: a cross-sectional study. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 424	8.3	32
145	Alcohol consumption and breast cancer risk among women in three sub-Saharan African countries. <i>PLoS ONE</i> , <b>2014</b> , 9, e106908	3.7	32
144	Inhibition of Copper Transport Induces Apoptosis in Triple-Negative Breast Cancer Cells and Suppresses Tumor Angiogenesis. <i>Molecular Cancer Therapeutics</i> , <b>2019</b> , 18, 873-885	6.1	31
143	Differentiation and loss of malignant character of spontaneous pulmonary metastases in patient-derived breast cancer models. <i>Cancer Research</i> , <b>2014</b> , 74, 7406-17	10.1	30
142	A functionally significant SNP in TP53 and breast cancer risk in African-American women. <i>Npj Breast Cancer</i> , <b>2017</b> , 3, 5	7.8	29
141	Association of Metformin Use with Outcomes in Advanced Endometrial Cancer Treated with Chemotherapy. <i>PLoS ONE</i> , <b>2016</b> , 11, e0147145	3.7	29
140	Building local capacity for genomics research in Africa: recommendations from analysis of publications in Sub-Saharan Africa from 2004 to 2013. <i>Global Health Action</i> , <b>2016</b> , 9, 31026	3	29
139	A comprehensive examination of breast cancer risk loci in African American women. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 5518-26	5.6	28
138	Characterization of the Cancer Spectrum in Men With Germline BRCA1 and BRCA2 Pathogenic Variants: Results From the Consortium of Investigators of Modifiers of BRCA1/2 (CIMBA). <i>JAMA Oncology</i> , <b>2020</b> , 6, 1218-1230	13.4	25
137	Randomized Noninferiority Trial of Telephone vs In-Person Disclosure of Germline Cancer Genetic Test Results. <i>Journal of the National Cancer Institute</i> , <b>2018</b> , 110, 985-993	9.7	25
136	Inheritance of deleterious mutations at both BRCA1 and BRCA2 in an international sample of 32,295 women. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 112	8.3	25
135	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 64	8.3	25

134	Evaluation of the Quality of Adjuvant Endocrine Therapy Delivery for Breast Cancer Care in the United States. <i>JAMA Oncology</i> , <b>2017</b> , 3, 928-935	13.4	24
133	Genetic variants in microRNA and microRNA biogenesis pathway genes and breast cancer risk among women of African ancestry. <i>Human Genetics</i> , <b>2016</b> , 135, 1145-59	6.3	24
132	Intensive Surveillance with Biannual Dynamic Contrast-Enhanced Magnetic Resonance Imaging Downstages Breast Cancer in Mutation Carriers. <i>Clinical Cancer Research</i> , <b>2019</b> , 25, 1786-1794	12.9	23
131	Identification of novel common breast cancer risk variants at the 6q25 locus mong Latinas. <i>Breast Cancer Research</i> , <b>2019</b> , 21, 3	8.3	23
130	Genetic variants demonstrating flip-flop phenomenon and breast cancer risk prediction among women of African ancestry. <i>Breast Cancer Research and Treatment</i> , <b>2018</b> , 168, 703-712	4.4	22
129	Association of Genomic Domains in and with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , <b>2020</b> , 80, 624-638	10.1	22
128	Prevalence of Inherited Mutations in Breast Cancer Predisposition Genes among Women in Uganda and Cameroon. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 359-367	4	22
127	Height and Body Mass Index as Modifiers of Breast Cancer Risk in BRCA1/2 Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 350-364	9.7	22
126	Candidate genetic modifiers for breast and ovarian cancer risk in BRCA1 and BRCA2 mutation carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 308-16	4	20
125	Closing the Cancer Divide Through Ubuntu: Information and Communication Technology-Powered Models for Global Radiation Oncology. <i>International Journal of Radiation Oncology Biology Physics</i> , <b>2016</b> , 94, 440-9	4	20
124	Efficacy of Anti-HER2 Agents in Combination With Adjuvant or Neoadjuvant Chemotherapy for Early and Locally Advanced HER2-Positive Breast Cancer Patients: A Network Meta-Analysis. <i>Frontiers in Oncology</i> , <b>2018</b> , 8, 156	5.3	20
123	Gain-of-Function Mutant p53 R273H Interacts with Replicating DNA and PARP1 in Breast Cancer. <i>Cancer Research</i> , <b>2020</b> , 80, 394-405	10.1	20
122	Epidemiology of Triple-Negative Breast Cancer: A Review. Cancer Journal (Sudbury, Mass), 2021, 27, 8-1	б <u>е</u> .2	20
121	Physical activity during adolescence and young adulthood and the risk of breast cancer in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , <b>2018</b> , 169, 561-571	4.4	19
120	CDKN2 gene deletion is not found in chronic lymphoid leukaemias of B- and T-cell origin but is frequent in acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , <b>1995</b> , 91, 865-70	4.5	19
119	Genetic and Epigenetic Regulation of TOX3 Expression in Breast Cancer. <i>PLoS ONE</i> , <b>2016</b> , 11, e0165559	3.7	19
118	Afatinib efficacy against squamous cell carcinoma of the head and neck cell lines in vitro and in vivo. <i>Targeted Oncology</i> , <b>2015</b> , 10, 501-8	5	18
117	Heterogeneity in hormone-receptor status and survival outcomes among women with synchronous and metachronous bilateral breast cancers. <i>Breast</i> , <b>2015</b> , 24, 131-6	3.6	18

116	Disparities in cancer care: a worldwide perspective and roadmap for change. <i>Journal of Clinical Oncology</i> , <b>2006</b> , 24, 2135-6	2.2	18
115	Community clinical practice patterns and mortality in patients with intermediate oncotype DX recurrence scores: Who benefits from chemotherapy?. <i>Cancer</i> , <b>2019</b> , 125, 213-222	6.4	18
114	Pilot study demonstrating potential association between breast cancer image-based risk phenotypes and genomic biomarkers. <i>Medical Physics</i> , <b>2014</b> , 41, 031917	4.4	17
113	Relationships between computer-extracted mammographic texture pattern features and. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 424	8.3	17
112	BRCA1 and BRCA2 pathogenic sequence variants in women of African origin or ancestry. <i>Human Mutation</i> , <b>2019</b> , 40, 1781-1796	4.7	16
111	Implementing newborn screening for sickle cell disease as part of immunisation programmes in Nigeria: a feasibility study. <i>Lancet Haematology,the</i> , <b>2020</b> , 7, e534-e540	14.6	16
110	Pilot Survey of Breast Cancer Management in Sub-Saharan Africa. <i>Journal of Global Oncology</i> , <b>2017</b> , 3, 194-200	2.6	16
109	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 61	8.3	16
108	Molecular Subtype-Specific Expression of MicroRNA-29c in Breast Cancer Is Associated with CpG Dinucleotide Methylation of the Promoter. <i>PLoS ONE</i> , <b>2015</b> , 10, e0142224	3.7	16
107	LncRNA BLAT1 is Upregulated in Basal-like Breast Cancer through Epigenetic Modifications. <i>Scientific Reports</i> , <b>2018</b> , 8, 15572	4.9	16
106	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , <b>2016</b> , 141, 386-401	4.9	15
105	Loss of BRCA1 in the Cells of Origin of Ovarian Cancer Induces Glycolysis: A Window of Opportunity for Ovarian Cancer Chemoprevention. <i>Cancer Prevention Research</i> , <b>2017</b> , 10, 255-266	3.2	15
104	Household air pollution and chronic hypoxia in the placenta of pregnant Nigerian women: A randomized controlled ethanol Cookstove intervention. <i>Science of the Total Environment</i> , <b>2018</b> , 619-620, 212-220	10.2	15
103	Population-dependent Intron Retention and DNA Methylation in Breast Cancer. <i>Molecular Cancer Research</i> , <b>2018</b> , 16, 461-469	6.6	15
102	Mutations in context: implications of BRCA testing in diverse populations. Familial Cancer, 2018, 17, 47	1-3483	15
101	Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , <b>2017</b> , 161, 117-134	4.4	15
100	The spectrum of BRCA1 and BRCA2 pathogenic sequence variants in Middle Eastern, North African, and South European countries. <i>Human Mutation</i> , <b>2019</b> , 40, e1-e23	4.7	14
99	Oestrogen receptor status and survival in women with BRCA2-associated breast cancer. <i>British Journal of Cancer</i> , <b>2019</b> , 120, 398-403	8.7	13

# (2020-2019)

98	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , <b>2019</b> , 121, 180-192	8.7	13
97	Clinical interpretation of pathogenic ATM and CHEK2 variants on multigene panel tests: navigating moderate risk. <i>Familial Cancer</i> , <b>2018</b> , 17, 495-505	3	13
96	Germline Rare Coding Variants and Risk of Pancreatic Cancer in Minority Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2018</b> , 27, 1364-1370	4	13
95	Susceptibility gene for familial acute myeloid leukemia associated with loss of 5q and/or 7q is not localized on the commonly deleted portion of 5q <b>2000</b> , 28, 164-172		13
94	Household air pollution and angiogenic factors in pregnant Nigerian women: A randomized controlled ethanol cookstove intervention. <i>Science of the Total Environment</i> , <b>2017</b> , 599-600, 2175-2181	10.2	12
93	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 1016-1026	4	12
92	Intrinsic adriamycin resistance in p53-mutated breast cancer is related to the miR-30c/FANCF/REV1-mediated DNA damage response. <i>Cell Death and Disease</i> , <b>2019</b> , 10, 666	9.8	12
91	The :p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , <b>2019</b> , 5, 38	7.8	12
90	Development of a Breast Cancer Risk Prediction Model for Women in Nigeria. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2018</b> , 27, 636-643	4	11
89	An Evaluation of Psychosocial and Religious Belief Differences in a Diverse Racial and Socioeconomic Urban Cancer Population. <i>Journal of Racial and Ethnic Health Disparities</i> , <b>2017</b> , 4, 140-14	.8 <sup>3.5</sup>	10
88	Infrastructural Challenges Lead to Delay of Curative Radiotherapy in Nigeria. <i>JCO Global Oncology</i> , <b>2020</b> , 6, 269-276	3.7	10
87	The impact of site-specific digital histology signatures on deep learning model accuracy and bias. <i>Nature Communications</i> , <b>2021</b> , 12, 4423	17.4	10
86	Propensity score analysis of the prognostic value of genomic assays for breast cancer in diverse populations using the National Cancer Data Base. <i>Cancer</i> , <b>2020</b> , 126, 4013-4022	6.4	9
85	Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic Epidemiology</i> , <b>2020</b> , 44, 442-468	2.6	9
84	Genetic anticipation in BRCA1/BRCA2 families after controlling for ascertainment bias and cohort effect. <i>Cancer</i> , <b>2016</b> , 122, 1913-20	6.4	9
83	Evaluating Polygenic Risk Scores for Breast Cancer in Women of African Ancestry. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 1168-1176	9.7	9
82	Breast Cancer Risk-Associated SNPs in the Promoter Form KLF5- and ZEB1-Binding Sites that Influence the Cellular Response to Paclitaxel. <i>Molecular Cancer Research</i> , <b>2019</b> , 17, 2244-2256	6.6	8
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