

# Britta Weigelt

## List of Publications by Year in descending order

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244  
papers

21,060  
citations

9775

73  
h-index

11928

134  
g-index

250  
all docs

250  
docs citations

250  
times ranked

27181  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic characterization of small cell carcinomas of the uterine cervix. <i>Molecular Oncology</i> , 2022, 16, 833-845.	2.1	14
2	INK4 Tumor Suppressor Proteins Mediate Resistance to CDK4/6 Kinase Inhibitors. <i>Cancer Discovery</i> , 2022, 12, 356-371.	7.7	68
3	The clinical behavior and genomic features of the so-called adenoid cystic carcinomas of the solid variant with basaloid features. <i>Modern Pathology</i> , 2022, 35, 193-201.	2.9	25
4	TSC2-mutant uterine sarcomas with JAZF1-SUZ12 fusions demonstrate hybrid features of endometrial stromal sarcoma and PEComa and are responsive to mTOR inhibition. <i>Modern Pathology</i> , 2022, 35, 117-127.	2.9	16
5	Morphologic and Genomic Characteristics of Breast Cancers Occurring in Individuals with Lynch Syndrome. <i>Clinical Cancer Research</i> , 2022, 28, 404-413.	3.2	13
6	Stromal <i>MED12</i> exon 2 mutations in complex fibroadenomas of the breast. <i>Journal of Clinical Pathology</i> , 2022, 75, 133-136.	1.0	2
7	Pathogenesis of Triple-Negative Breast Cancer. <i>Annual Review of Pathology: Mechanisms of Disease</i> , 2022, 17, 181-204.	9.6	132
8	<i>ATM</i> Germline-Mutated Gastroesophageal Junction Adenocarcinomas: Clinical Descriptors, Molecular Characteristics, and Potential Therapeutic Implications. <i>Journal of the National Cancer Institute</i> , 2022, 114, 761-770.	3.0	3
9	Genomic characterization of metastatic patterns from prospective clinical sequencing of 25,000 patients. <i>Cell</i> , 2022, 185, 563-575.e11.	13.5	223
10	Intratumor genetic heterogeneity and clonal evolution to decode endometrial cancer progression. <i>Oncogene</i> , 2022, 41, 1835-1850.	2.6	9
11	Hyperthermic intraperitoneal chemotherapy (HIPEC) with carboplatin induces distinct transcriptomic changes in ovarian tumor and normal tissues. <i>Gynecologic Oncology</i> , 2022, 165, 239-247.	0.6	9
12	Same-Cell Co-Occurrence of RAS Hotspot and BRAF V600E Mutations in Treatment-Naive Colorectal Cancer. <i>JCO Precision Oncology</i> , 2022, 6, e2100365.	1.5	1
13	Genomic landscape of endometrial carcinomas of no specific molecular profile. <i>Modern Pathology</i> , 2022, 35, 1269-1278.	2.9	33
14	Histopathologic features and molecular genetic landscape of HER2-amplified endometrial carcinomas. <i>Modern Pathology</i> , 2022, 35, 962-971.	2.9	22
15	Recurrent <i>WWTR1</i> <i>S89W</i> mutations and Hippo pathway deregulation in clear cell carcinomas of the cervix. <i>Journal of Pathology</i> , 2022, 257, 635-649.	2.1	2
16	Clinical-pathologic characteristics and response to neoadjuvant chemotherapy in triple-negative low Ki-67 proliferation (TNLP) breast cancers. <i>Npj Breast Cancer</i> , 2022, 8, 51.	2.3	9
17	MAPK Pathway Genetic Alterations Are Associated with Prolonged Overall Survival in Low-Grade Serous Ovarian Carcinoma. <i>Clinical Cancer Research</i> , 2022, 28, 4456-4465.	3.2	25
18	Evaluation of TERT mRNA expression using RNAscope®: A potential histopathologic diagnostic and prognostic tool. <i>Pathology Research and Practice</i> , 2022, 233, 153892.	1.0	2

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19	Immunogenicity and therapeutic targeting of a public neoantigen derived from mutated PIK3CA. <i>Nature Medicine</i> , 2022, 28, 946-957.	15.2	50
20	Fundamental immune oncogenicity trade-offs define driver mutation fitness. <i>Nature</i> , 2022, 606, 172-179.	13.7	23
21	Treatment of ovarian clear cell carcinoma with immune checkpoint blockade: a case series. <i>International Journal of Gynecological Cancer</i> , 2022, , ijgc-2022-003430.	1.2	5
22	Molecular Subclasses of Clear Cell Ovarian Carcinoma and Their Impact on Disease Behavior and Outcomes. <i>Clinical Cancer Research</i> , 2022, 28, 4947-4956.	3.2	22
23	Pathogenic ATM Mutations in Cancer and a Genetic Basis for Radiotherapeutic Efficacy. <i>Journal of the National Cancer Institute</i> , 2021, 113, 266-273.	3.0	38
24	Problematic breast tumors reassessed in light of novel molecular data. <i>Modern Pathology</i> , 2021, 34, 38-47.	2.9	25
25	The genetic landscape of metaplastic breast cancers and uterine carcinosarcomas. <i>Molecular Oncology</i> , 2021, 15, 1024-1039.	2.1	21
26	Whole exome sequencing analysis of juvenile papillomatosis and coexisting breast carcinoma. <i>Journal of Pathology: Clinical Research</i> , 2021, 7, 113-120.	1.3	4
27	Massively parallel sequencing analysis of 68 gastric-type cervical adenocarcinomas reveals mutations in cell cycle-related genes and potentially targetable mutations. <i>Modern Pathology</i> , 2021, 34, 1213-1225.	2.9	28
28	Clonal relationship and directionality of progression of synchronous endometrial and ovarian carcinomas in patients with DNA mismatch repair-deficiency associated syndromes. <i>Modern Pathology</i> , 2021, 34, 994-1007.	2.9	19
29	Metaplastic carcinomas of the breast without evidence of epithelial differentiation: a diagnostic approach for management. <i>Histopathology</i> , 2021, 78, 759-771.	1.6	13
30	Precision Radiotherapy: Reduction in Radiation for Oropharyngeal Cancer in the 30 ROC Trial. <i>Journal of the National Cancer Institute</i> , 2021, 113, 742-751.	3.0	98
31	Ultraviolet radiation drives mutations in a subset of mucosal melanomas. <i>Nature Communications</i> , 2021, 12, 259.	5.8	27
32	Interobserver Variation of PD-L1 SP142 Immunohistochemistry Interpretation in Breast Carcinoma: A Study of 79 Cases Using Whole Slide Imaging. <i>Archives of Pathology and Laboratory Medicine</i> , 2021, 145, 1132-1137.	1.2	11
33	Activation of the IFN Signaling Pathway is Associated with Resistance to CDK4/6 Inhibitors and Immune Checkpoint Activation in ER-Positive Breast Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 4870-4882.	3.2	49
34	Clinicopathologic and Genomic Analysis of TP53-Mutated Endometrial Carcinomas. <i>Clinical Cancer Research</i> , 2021, 27, 2613-2623.	3.2	49
35	Targeting galectin-3 with a high-affinity antibody for inhibition of high-grade serous ovarian cancer and other MUC16/CA-125-expressing malignancies. <i>Scientific Reports</i> , 2021, 11, 3718.	1.6	18
36	Homologous recombination deficiency: how genomic signatures are generated. <i>Current Opinion in Genetics and Development</i> , 2021, 66, 93-100.	1.5	13

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37	Genomic Alterations in <i>PIK3CA</i> -Mutated Breast Cancer Result in mTORC1 Activation and Limit the Sensitivity to PI3K± Inhibitors. <i>Cancer Research</i> , 2021, 81, 2470-2480.	0.4	20
38	Mesonephric and mesonephric-like carcinomas of the female genital tract: molecular characterization including cases with mixed histology and matched metastases. <i>Modern Pathology</i> , 2021, 34, 1570-1587.	2.9	57
39	TERT promoter hotspot mutations and gene amplification in metaplastic breast cancer. <i>Npj Breast Cancer</i> , 2021, 7, 43.	2.3	16
40	Genetic interactions among Brca1, Brca2, Palb2, and Trp53 in mammary tumor development. <i>Npj Breast Cancer</i> , 2021, 7, 45.	2.3	7
41	Genetic characterisation of adult primary pleomorphic uterine rhabdomyosarcoma and comparison with uterine carcinosarcoma. <i>Histopathology</i> , 2021, 79, 176-186.	1.6	4
42	Independent real-world application of a clinical-grade automated prostate cancer detection system. <i>Journal of Pathology</i> , 2021, 254, 147-158.	2.1	57
43	Genetic and molecular subtype heterogeneity in newly diagnosed early- and advanced-stage endometrial cancer. <i>Gynecologic Oncology</i> , 2021, 161, 535-544.	0.6	16
44	Molecular characterization of high-grade serous ovarian cancers occurring in younger and older women. <i>Gynecologic Oncology</i> , 2021, 161, 545-552.	0.6	8
45	The genomic landscape of carcinomas with mucinous differentiation. <i>Scientific Reports</i> , 2021, 11, 9478.	1.6	9
46	Histologic and genomic features of breast cancers with alterations affecting the SWI/SNF (SMARC) genes. <i>Modern Pathology</i> , 2021, 34, 1850-1859.	2.9	3
47	Poor response to neoadjuvant chemotherapy in metaplastic breast carcinoma. <i>Npj Breast Cancer</i> , 2021, 7, 96.	2.3	38
48	Paired Tumor-Normal Sequencing Provides Insights into TP53-Related Cancer Spectrum in Li-Fraumeni Patients. <i>Journal of the National Cancer Institute</i> , 2021, , .	3.0	6
49	Pattern of disease and response to pembrolizumab in recurrent cervical cancer. <i>Gynecologic Oncology Reports</i> , 2021, 37, 100831.	0.3	4
50	Diagnosis and management of an endometrial cancer patient with Cowden syndrome. <i>Gynecologic Oncology</i> , 2021, 163, 14-21.	0.6	6
51	Recurrence biomarkers of triple negative breast cancer treated with neoadjuvant chemotherapy and anti-EGFR antibodies. <i>Npj Breast Cancer</i> , 2021, 7, 124.	2.3	7
52	Clinical utility of next-generation sequencing-based ctDNA testing for common and novel ALK fusions. <i>Lung Cancer</i> , 2021, 159, 66-73.	0.9	17
53	Spectrum of <i>BRAF</i> Mutations and Gene Rearrangements in Ovarian Serous Carcinoma. <i>JCO Precision Oncology</i> , 2021, 5, 1480-1492.	1.5	8
54	Ki67 Assessment in Breast Cancer: Are We There Yet?. <i>Journal of the National Cancer Institute</i> , 2021, 113, 797-798.	3.0	7

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55	Genomic Profiling Aids Classification of Diagnostically Challenging Uterine Mesenchymal Tumors With Myomelanocytic Differentiation. <i>American Journal of Surgical Pathology</i> , 2021, 45, 77-92.	2.1	30
56	AKT1 E17K Inhibits Cancer Cell Migration by Abrogating $\beta$ -Catenin Signaling. <i>Molecular Cancer Research</i> , 2021, 19, 573-584.	1.5	10
57	Metastasis and Immune Evasion from Extracellular cGAMP Hydrolysis. <i>Cancer Discovery</i> , 2021, 11, 1212-1227.	7.7	139
58	Germline RAD51B variants confer susceptibility to breast and ovarian cancers deficient in homologous recombination. <i>Npj Breast Cancer</i> , 2021, 7, 135.	2.3	9
59	Diverse alterations associated with resistance to KRAS(G12C) inhibition. <i>Nature</i> , 2021, 599, 679-683.	13.7	183
60	Geometric network analysis provides prognostic information in patients with high grade serous carcinoma of the ovary treated with immune checkpoint inhibitors. <i>Npj Genomic Medicine</i> , 2021, 6, 99.	1.7	13
61	Pancreatoblastomas and mixed and pure acinar cell carcinomas share epigenetic signatures distinct from other neoplasms of the pancreas. <i>Modern Pathology</i> , 2021, , .	2.9	3
62	ESR1 hotspot mutations in endometrial stromal sarcoma with high-grade transformation and endocrine treatment. <i>Modern Pathology</i> , 2021, , .	2.9	5
63	HER2-Enriched Subtype and ERBB2 Expression in HER2-Positive Breast Cancer Treated with Dual HER2 Blockade. <i>Journal of the National Cancer Institute</i> , 2020, 112, 46-54.	3.0	97
64	Sclerosing epithelioid mesenchymal neoplasm of the pancreas—A proposed new entity. <i>Modern Pathology</i> , 2020, 33, 456-467.	2.9	10
65	Histologic spectrum of polymorphous adenocarcinoma of the salivary gland harbor genetic alterations affecting PRKD genes. <i>Modern Pathology</i> , 2020, 33, 65-73.	2.9	29
66	Evaluating Clonal Hematopoiesis in Tumor-Infiltrating Leukocytes in Breast Cancer and Secondary Hematologic Malignancies. <i>Journal of the National Cancer Institute</i> , 2020, 112, 107-110.	3.0	10
67	Evaluation of the Predictive Role of Tumor Immune Infiltrate in Patients with HER2-Positive Breast Cancer Treated with Neoadjuvant Anti-HER2 Therapy without Chemotherapy. <i>Clinical Cancer Research</i> , 2020, 26, 738-745.	3.2	31
68	Immunohistochemical assessment of HRASQ61R mutations in breast adenomyoepitheliomas. <i>Histopathology</i> , 2020, 76, 865-874.	1.6	19
69	Identification of recurrent FHL2-GLI2 oncogenic fusion in sclerosing stromal tumors of the ovary. <i>Nature Communications</i> , 2020, 11, 44.	5.8	34
70	Hormone receptor and HER2 assessment in breast carcinoma metastatic to bone: A comparison between FNA cell blocks and decalcified core needle biopsies. <i>Cancer Cytopathology</i> , 2020, 128, 133-145.	1.4	4
71	Clinical outcomes of patients with POLE mutated endometrioid endometrial cancer. <i>Gynecologic Oncology</i> , 2020, 156, 194-202.	0.6	35
72	Immunohistochemical analysis of IDH2 R172 hotspot mutations in breast papillary neoplasms: applications in the diagnosis of tall cell carcinoma with reverse polarity. <i>Modern Pathology</i> , 2020, 33, 1056-1064.	2.9	35

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73	Distinct Classes of Complex Structural Variation Uncovered across Thousands of Cancer Genome Graphs. <i>Cell</i> , 2020, 183, 197-210.e32.	13.5	141
74	Acquisition of APOBEC Mutagenesis and Microsatellite Instability Signatures in the Development of Brain Metastases in Low-Grade, Early-Stage Endometrioid Endometrial Carcinoma. <i>JCO Precision Oncology</i> , 2020, 4, 1217-1223.	1.5	1
75	The genomic landscape of metastatic histologic special types of invasive breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 53.	2.3	27
76	Machine learning-based prediction of microsatellite instability and high tumor mutation burden from contrast-enhanced computed tomography in endometrial cancers. <i>Scientific Reports</i> , 2020, 10, 17769.	1.6	35
77	Whole-exome analysis of metaplastic breast carcinomas with extensive osseous differentiation. <i>Histopathology</i> , 2020, 77, 321-326.	1.6	7
78	Mutations in BRCA1 and BRCA2 differentially affect the tumor microenvironment and response to checkpoint blockade immunotherapy. <i>Nature Cancer</i> , 2020, 1, 1188-1203.	5.7	114
79	A prospective multicenter international single-arm observational study on the oncological safety of the sentinel lymph node algorithm in stage I intermediate-risk endometrial cancer (SELECT, SEntinel) <i>Tj ETQq1 1 0.784314 rgBT /Over</i> 1627-1632.	1.2	16
80	FOXA1 Mutations Reveal Distinct Chromatin Profiles and Influence Therapeutic Response in Breast Cancer. <i>Cancer Cell</i> , 2020, 38, 534-550.e9.	7.7	67
81	Neuroendocrine tumours of the breast: a genomic comparison with mucinous breast cancers and neuroendocrine tumours of other anatomic sites. <i>Journal of Clinical Pathology</i> , 2020, , jclinpath-2020-207052.	1.0	5
82	Histologic Classification and Molecular Signature of Polymorphous Adenocarcinoma (PAC) and Cribriform Adenocarcinoma of Salivary Gland (CASG). <i>American Journal of Surgical Pathology</i> , 2020, 44, 545-552.	2.1	39
83	Unraveling tumor-immune heterogeneity in advanced ovarian cancer uncovers immunogenic effect of chemotherapy. <i>Nature Genetics</i> , 2020, 52, 582-593.	9.4	136
84	High-grade transformation of low-grade endometrial stromal sarcomas lacking YWHAE and BCOR genetic abnormalities. <i>Modern Pathology</i> , 2020, 33, 1861-1870.	2.9	26
85	Oncogenic properties and signaling basis of the PAX8-GLIS3 fusion gene. <i>International Journal of Cancer</i> , 2020, 147, 2253-2264.	2.3	10
86	BRCA Mutations, Homologous DNA Repair Deficiency, Tumor Mutational Burden, and Response to Immune Checkpoint Inhibition in Recurrent Ovarian Cancer. <i>JCO Precision Oncology</i> , 2020, 4, 665-679.	1.5	29
87	Pleomorphic adenomas and mucoepidermoid carcinomas of the breast are underpinned by fusion genes. <i>Npj Breast Cancer</i> , 2020, 6, 20.	2.3	25
88	Genomic profiling of primary and recurrent adult granulosa cell tumors of the ovary. <i>Modern Pathology</i> , 2020, 33, 1606-1617.	2.9	38
89	Clinical and pathologic features associated with PD-L1 (SP142) expression in stromal tumor-infiltrating immune cells of triple-negative breast carcinoma. <i>Modern Pathology</i> , 2020, 33, 2221-2232.	2.9	23
90	Mutant FOXL2C134W Hijacks SMAD4 and SMAD2/3 to Drive Adult Granulosa Cell Tumors. <i>Cancer Research</i> , 2020, 80, 3466-3479.	0.4	29

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91	A P53-Independent DNA Damage Response Suppresses Oncogenic Proliferation and Genome Instability. <i>Cell Reports</i> , 2020, 30, 1385-1399.e7.	2.9	29
92	ARID1A determines luminal identity and therapeutic response in estrogen-receptor-positive breast cancer. <i>Nature Genetics</i> , 2020, 52, 198-207.	9.4	140
93	Alterations in PTEN and ESR1 promote clinical resistance to alpelisib plus aromatase inhibitors. <i>Nature Cancer</i> , 2020, 1, 382-393.	5.7	96
94	Endometrial Carcinomas with a "Serous" Component in Young Women Are Enriched for DNA Mismatch Repair Deficiency, Lynch Syndrome, and POLE Exonuclease Domain Mutations. <i>American Journal of Surgical Pathology</i> , 2020, 44, 641-648.	2.1	34
95	Whole-Exome Sequencing Analysis of the Progression from Non-"Low-Grade Ductal Carcinoma In Situ" to Invasive Ductal Carcinoma. <i>Clinical Cancer Research</i> , 2020, 26, 3682-3693.	3.2	42
96	HER2-Mediated Internalization of Cytotoxic Agents in ERBB2 Amplified or Mutant Lung Cancers. <i>Cancer Discovery</i> , 2020, 10, 674-687.	7.7	149
97	The impact of poly ADP ribose polymerase (PARP) inhibitors on clonal hematopoiesis.. <i>Journal of Clinical Oncology</i> , 2020, 38, 1513-1513.	0.8	16
98	Gastric-type adenocarcinoma of the cervix: Genomic drivers and clinical outcomes.. <i>Journal of Clinical Oncology</i> , 2020, 38, 6030-6030.	0.8	1
99	Homologous recombination DNA repair defects in PALB2-associated breast cancers. <i>Npj Breast Cancer</i> , 2019, 5, 23.	2.3	39
100	How Did We Get There? The Progression from Ductal Carcinoma In Situ to Invasive Ductal Carcinoma. <i>Current Breast Cancer Reports</i> , 2019, 11, 175-184.	0.5	0
101	Molecular profiling and molecular classification of endometrioid ovarian carcinomas. <i>Gynecologic Oncology</i> , 2019, 154, 516-523.	0.6	62
102	Whole-exome sequencing and RNA sequencing analyses of acinic cell carcinomas of the breast. <i>Histopathology</i> , 2019, 75, 931-937.	1.6	16
103	PAX8-"GLIS3 gene fusion is a pathognomonic genetic alteration of hyalinizing trabecular tumors of the thyroid. <i>Modern Pathology</i> , 2019, 32, 1734-1743.	2.9	38
104	Targeting the Mevalonate Pathway to Overcome Acquired Anti-HER2 Treatment Resistance in Breast Cancer. <i>Molecular Cancer Research</i> , 2019, 17, 2318-2330.	1.5	41
105	Assessment of HMGA2 and PLAG1 rearrangements in breast adenomyoepitheliomas. <i>Npj Breast Cancer</i> , 2019, 5, 6.	2.3	21
106	The Landscape of Somatic Genetic Alterations in Breast Cancers from CHEK2 Germline Mutation Carriers. <i>JNCI Cancer Spectrum</i> , 2019, 3, pkz027.	1.4	20
107	V211D Mutation in MEK1 Causes Resistance to MEK Inhibitors in Colon Cancer. <i>Cancer Discovery</i> , 2019, 9, 1182-1191.	7.7	27
108	Secretory carcinoma of the breast: clinicopathologic profile of 14 cases emphasising distant metastatic potential. <i>Histopathology</i> , 2019, 75, 213-224.	1.6	46

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109	Genomic analysis of recurrences and high-grade forms of polymorphous adenocarcinoma. <i>Histopathology</i> , 2019, 75, 193-201.	1.6	10
110	Micropapillary variant of mucinous carcinoma of the breast shows genetic alterations intermediate between those of mucinous carcinoma and micropapillary carcinoma. <i>Histopathology</i> , 2019, 75, 139-145.	1.6	22
111	Solid pseudopapillary neoplasms of the pancreas are dependent on the Wnt pathway. <i>Molecular Oncology</i> , 2019, 13, 1684-1692.	2.1	21
112	The repertoire of genetic alterations in salivary duct carcinoma including a novel HNRNP3-ALK rearrangement. <i>Human Pathology</i> , 2019, 88, 66-77.	1.1	38
113	Massively parallel sequencing analysis of benign melanocytic naevi. <i>Histopathology</i> , 2019, 75, 29-38.	1.6	12
114	Functional and topographic effects on DNA methylation in IDH1/2 mutant cancers. <i>Scientific Reports</i> , 2019, 9, 16830.	1.6	29
115	Radiogenomics Analysis of Intratumor Heterogeneity in a Patient With High-Grade Serous Ovarian Cancer. <i>JCO Precision Oncology</i> , 2019, 3, 1-9.	1.5	10
116	FOXA1 upregulation promotes enhancer and transcriptional reprogramming in endocrine-resistant breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 26823-26834.	3.3	103
117	High-intensity sequencing reveals the sources of plasma circulating cell-free DNA variants. <i>Nature Medicine</i> , 2019, 25, 1928-1937.	15.2	485
118	Endometrial Cancers in <i>BRCA1</i> or <i>BRCA2</i> Germline Mutation Carriers: Assessment of Homologous Recombination DNA Repair Defects. <i>JCO Precision Oncology</i> , 2019, 3, 1-11.	1.5	19
119	The role of a monoclonal antibody 11C8B1 as a diagnostic marker of IDH2-mutated sinonasal undifferentiated carcinoma. <i>Modern Pathology</i> , 2019, 32, 205-215.	2.9	22
120	Lobular Carcinomas <i>In Situ</i> Display Intralesion Genetic Heterogeneity and Clonal Evolution in the Progression to Invasive Lobular Carcinoma. <i>Clinical Cancer Research</i> , 2019, 25, 674-686.	3.2	44
121	The clinical use of circulating tumor cells (CTCs) enumeration for staging of metastatic breast cancer (MBC): International expert consensus paper. <i>Critical Reviews in Oncology/Hematology</i> , 2019, 134, 39-45.	2.0	200
122	Somatic genetic alterations in synchronous and metachronous low-grade serous tumours and high-grade carcinomas of the adnexa. <i>Histopathology</i> , 2019, 74, 638-650.	1.6	11
123	Analysis of mutational signatures in primary and metastatic endometrial cancer reveals distinct patterns of DNA repair defects and shifts during tumor progression. <i>Gynecologic Oncology</i> , 2019, 152, 11-19.	0.6	66
124	Recurrent <i>MED12</i> exon 2 mutations in benign breast fibroepithelial lesions in adolescents and young adults. <i>Journal of Clinical Pathology</i> , 2019, 72, 258-262.	1.0	22
125	The Genomic Landscape of Mucinous Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 737-741.	3.0	68
126	Genomic Applications in Gynecologic Malignancies. , 2019, , 445-469.		0



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127	Genetic hallmarks of recurrent/metastatic adenoid cystic carcinoma. <i>Journal of Clinical Investigation</i> , 2019, 129, 4276-4289.	3.9	134
128	The molecular genetic make-up of male breast cancer. <i>Endocrine-Related Cancer</i> , 2019, 26, 779-794.	1.6	27
129	The Landscape of Somatic Genetic Alterations in Breast Cancers From ATM Germline Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2018, 110, 1030-1034.	3.0	90
130	Mutation Profiling of Key Cancer Genes in Primary Breast Cancers and Their Distant Metastases. <i>Cancer Research</i> , 2018, 78, 3112-3121.	0.4	57
131	Solid papillary breast carcinomas resembling the tall cell variant of papillary thyroid neoplasms (solid papillary carcinomas with reverse polarity) harbour recurrent mutations affecting <i>IDH2</i> and <i>PIK3CA</i> : a validation cohort. <i>Histopathology</i> , 2018, 73, 339-344.	1.6	44
132	The value of cell-free DNA for molecular pathology. <i>Journal of Pathology</i> , 2018, 244, 616-627.	2.1	91
133	Triple-negative breast cancers – a panoply of cancer types. <i>Nature Reviews Clinical Oncology</i> , 2018, 15, 347-348.	12.5	45
134	Contralateral breast cancers: Independent cancers or metastases?. <i>International Journal of Cancer</i> , 2018, 142, 347-356.	2.3	37
135	Invasion in breast lesions: the role of the epithelial-stroma barrier. <i>Histopathology</i> , 2018, 72, 1075-1083.	1.6	25
136	Low PTEN levels and PIK3CA mutations predict resistance to neoadjuvant lapatinib and trastuzumab without chemotherapy in patients with HER2 over-expressing breast cancer. <i>Breast Cancer Research and Treatment</i> , 2018, 167, 731-740.	1.1	71
137	<i>MYBL1</i> rearrangements and <i>MYB</i> amplification in breast adenoid cystic carcinomas lacking the <i>MYB</i> - <i>NFIB</i> fusion gene. <i>Journal of Pathology</i> , 2018, 244, 143-150.	2.1	74
138	Reliability of Whole-Exome Sequencing for Assessing Intratumor Genetic Heterogeneity. <i>Cell Reports</i> , 2018, 25, 1446-1457.	2.9	76
139	Genetic heterogeneity and actionable mutations in HER2-positive primary breast cancers and their brain metastases. <i>Oncotarget</i> , 2018, 9, 20617-20630.	0.8	36
140	Loss of the FAT1 Tumor Suppressor Promotes Resistance to CDK4/6 Inhibitors via the Hippo Pathway. <i>Cancer Cell</i> , 2018, 34, 893-905.e8.	7.7	307
141	The Genomic Landscape of Endocrine-Resistant Advanced Breast Cancers. <i>Cancer Cell</i> , 2018, 34, 427-438.e6.	7.7	633
142	Loss-of-function mutations in ATP6AP1 and ATP6AP2 in granular cell tumors. <i>Nature Communications</i> , 2018, 9, 3533.	5.8	92
143	Massively parallel sequencing analysis of mucinous ovarian carcinomas: genomic profiling and differential diagnoses. <i>Gynecologic Oncology</i> , 2018, 150, 127-135.	0.6	41
144	Clinical Utility of Prospective Molecular Characterization in Advanced Endometrial Cancer. <i>Clinical Cancer Research</i> , 2018, 24, 5939-5947.	3.2	100

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145	Recurrent hotspot mutations in HRAS Q61 and PI3K-AKT pathway genes as drivers of breast adenomyoepitheliomas. <i>Nature Communications</i> , 2018, 9, 1816.	5.8	105
146	The clinical utility of prospective molecular characterization in advanced cervical and vulvovaginal cancer.. <i>Journal of Clinical Oncology</i> , 2018, 36, 5531-5531.	0.8	0
147	Breast Cancer Heterogeneity: Roles in Tumorigenesis and Therapeutic Implications. <i>Current Breast Cancer Reports</i> , 2017, 9, 34-44.	0.5	11
148	Genetic analysis of uterine adenosarcomas and phyllodes tumors of the breast. <i>Molecular Oncology</i> , 2017, 11, 913-926.	2.1	11
149	Whole-genome single-cell copy number profiling from formalin-fixed paraffin-embedded samples. <i>Nature Medicine</i> , 2017, 23, 376-385.	15.2	111
150	The Landscape of Somatic Genetic Alterations in Metaplastic Breast Carcinomas. <i>Clinical Cancer Research</i> , 2017, 23, 3859-3870.	3.2	129
151	Generation of conditional oncogenic chromosomal translocations using CRISPR-Cas9 genomic editing and homology-directed repair. <i>Journal of Pathology</i> , 2017, 242, 102-112.	2.1	23
152	Genetic analysis of a morphologically heterogeneous ovarian endometrioid carcinoma. <i>Histopathology</i> , 2017, 71, 480-487.	1.6	2
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