List of Publications by Year in descending order

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IVAN RIECHE

#	Article	IF	CITATIONS
1	Frequency of microsatellite instability (MSI) in upper tract urothelial carcinoma: comparison of the Bethesda panel and the Idylla MSI assay in a consecutively collected, multi-institutional cohort. Journal of Clinical Pathology, 2023, 76, 126-132.	2.0	7
2	Differential Expression of Genes Involved in Metabolism and Immune Response in Diffuse and Intestinal Gastric Cancers, a Pilot Ptudy. Biomedicines, 2022, 10, 240.	3.2	1
3	Value of the loss of heterozygosity to BRCA1 variant classification. Npj Breast Cancer, 2022, 8, 9.	5.2	2
4	Prognostic Value of Fusobacterium nucleatum after Abdominoperineal Resection for Anal Squamous Cell Carcinoma. Cancers, 2022, 14, 1606.	3.7	7
5	Dramatic In Vivo Efficacy of the EZH2-Inhibitor Tazemetostat in PBRM1-Mutated Human Chordoma Xenograft. Cancers, 2022, 14, 1486.	3.7	10
6	Upregulated flotillins and sphingosine kinase 2 derail AXL vesicular traffic to promote epithelial-mesenchymal transition. Journal of Cell Science, 2022, 135, .	2.0	6
7	PD-1 Blockade in Solid Tumors with Defects in Polymerase Epsilon. Cancer Discovery, 2022, 12, 1435-1448.	9.4	28
8	Human papilloma virus integration sites and genomic signatures in head and neck squamous cell carcinoma. Molecular Oncology, 2022, 16, 3001-3016.	4.6	7
9	Kindlinâ€1 modulates the EGFR pathway and predicts sensitivity to EGFR inhibitors across cancer types. Clinical and Translational Medicine, 2022, 12, e813.	4.0	0
10	BCG therapy downregulates HLA-I on malignant cells to subvert antitumor immune responses in bladder cancer. Journal of Clinical Investigation, 2022, 132, .	8.2	11
11	Comprehensive Genome Profiling in Patients With Metastatic Non–Small Cell Lung Cancer: The Precision Medicine Phase II Randomized SAFIR02-Lung/IFCT 1301 Trial. Clinical Cancer Research, 2022, 28, 4018-4026.	7.0	4
12	PARP inhibitors and radiation potentiate liver cell death in vitro. Do hepatocellular carcinomas have an achilles' heel?. Clinics and Research in Hepatology and Gastroenterology, 2021, 45, 101553.	1.5	11
13	Fineâ€needle aspiration as an alternative to core needle biopsy for tumour molecular profiling in precision oncology: prospective comparative study of nextâ€generation sequencing in cancer patients included in the SHIVA02 trial. Molecular Oncology, 2021, 15, 104-115.	4.6	10
14	Interleukin-8 Receptors CXCR1 and CXCR2 Are Not Expressed by Endothelial Colony-forming Cells. Stem Cell Reviews and Reports, 2021, 17, 628-638.	3.8	0
15	Human papilloma virus (HPV) integration signature in Cervical Cancer: identification of MACROD2 gene as HPV hot spot integration site. British Journal of Cancer, 2021, 124, 777-785.	6.4	44
16	Genomic Alterations in Head and Neck Squamous Cell Carcinoma: Level of Evidence According to ESMO Scale for Clinical Actionability of Molecular Targets (ESCAT). JCO Precision Oncology, 2021, 5, 215-226.	3.0	22
17	Pharmacologic Normalization of Pancreatic Cancer-Associated Fibroblast Secretome Impairs Prometastatic Cross-Talk With Macrophages. Cellular and Molecular Gastroenterology and Hepatology, 2021, 11, 1405-1436.	4.5	21
18	Hyperprogressive Disease After Pembrolizumab Treatment in Advanced Epstein-Barr Virus–Associated Gastric Adenocarcinoma With ERBB2 Amplification. JCO Precision Oncology, 2021, 5, 370-377.	3.0	2

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19	Molecular features of untreated breast cancer and initial metastatic event inform clinical decision-making and predict outcome: long-term results of ESOPE, a single-arm prospective multicenter study. Genome Medicine, 2021, 13, 44.	8.2	13
20	Prognostic value of intratumoral Fusobacterium nucleatum and association with immune-related gene expression in oral squamous cell carcinoma patients. Scientific Reports, 2021, 11, 7870.	3.3	31
21	High <i>in vitro</i> and <i>in vivo</i> synergistic activity between mTORC1 and PLK1 inhibition in adenocarcinoma NSCLC. Oncotarget, 2021, 12, 859-872.	1.8	4
22	Metastasis-suppressor NME1 controls the invasive switch of breast cancer by regulating MT1-MMP surface clearance. Oncogene, 2021, 40, 4019-4032.	5.9	19
23	Analysis of genomic and non-genomic signaling of estrogen receptor in PDX models of breast cancer treated with a combination of the PI3K inhibitor alpelisib (BYL719) and fulvestrant. Breast Cancer Research, 2021, 23, 57.	5.0	7
24	5′ Region Large Genomic Rearrangements in the BRCA1 Gene in French Families: Identification of a Tandem Triplication and Nine Distinct Deletions with Five Recurrent Breakpoints. Cancers, 2021, 13, 3171.	3.7	5
25	Definition of Biologically Distinct Groups of Conjunctival Melanomas According to Etiological Factors and Implications for Precision Medicine. Cancers, 2021, 13, 3836.	3.7	10
26	Circulating HPV DNA as a Marker for Early Detection of Relapse in Patients with Cervical Cancer. Clinical Cancer Research, 2021, 27, 5869-5877.	7.0	36
27	Phase I trial of copanlisib, a selective PI3K inhibitor, in combination with cetuximab in patients with recurrent and/or metastatic head and neck squamous cell carcinoma. Investigational New Drugs, 2021, 39, 1641-1648.	2.6	9
28	Spontaneous mouse lymphoma in patient-derived tumor xenografts: The importance of systematic analysis of xenografted human tumor tissues in preclinical efficacy trials. Translational Oncology, 2021, 14, 101133.	3.7	6
29	HRAS is a therapeutic target in malignant chemo-resistant adenomyoepithelioma of the breast. Journal of Hematology and Oncology, 2021, 14, 143.	17.0	7
30	The mitochondrially-localized nucleoside diphosphate kinase D (NME4) is a novel metastasis suppressor. BMC Biology, 2021, 19, 228.	3.8	21
31	Biopathological Significance of PIWI–piRNA Pathway Deregulation in Invasive Breast Carcinomas. Cancers, 2020, 12, 2833.	3.7	6
32	Genomic Instability Signature of Palindromic Non-Coding Somatic Mutations in Bladder Cancer. Cancers, 2020, 12, 2882.	3.7	13
33	Assessment of prognostic implication of a panel of oncogenes in bladder cancer and identification of a 3-gene signature associated with recurrence and progression risk in non-muscle-invasive bladder cancer. Scientific Reports, 2020, 10, 16641.	3.3	10
34	PLK1 inhibition exhibits strong anti-tumoral activity in CCND1-driven breast cancer metastases with acquired palbociclib resistance. Nature Communications, 2020, 11, 4053.	12.8	77
35	Human Aortic Valve Interstitial Cells Display Proangiogenic Properties During Calcific Aortic Valve Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2020, 41, 415-429.	2.4	12
36	Altered Expression of Three EGFR Posttranslational Regulators MDGI, MIG6, and EIG121 in Invasive Breast Carcinomas. Analytical Cellular Pathology, 2020, 2020, 1-10.	1.4	3

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37	EHD2 is a Predictive Biomarker of Chemotherapy Efficacy in Triple Negative Breast Carcinoma. Scientific Reports, 2020, 10, 7998.	3.3	5
38	MMP2 as an independent prognostic stratifier in oral cavity cancers. OncoImmunology, 2020, 9, 1754094.	4.6	15
39	BRCAness, SLFN11, and RB1 loss predict response to topoisomerase I inhibitors in triple-negative breast cancers. Science Translational Medicine, 2020, 12, .	12.4	86
40	A single droplet digital PCR for ESR1 activating mutations detection in plasma. Oncogene, 2020, 39, 2987-2995.	5.9	42
41	Tubulin polyglutamylation is a general traffic control mechanism in hippocampal neurons. Journal of Cell Science, 2020, 133, .	2.0	39
42	ZRANB2 and SYF2-mediated splicing programs converging on ECT2 are involved in breast cancer cell resistance to doxorubicin. Nucleic Acids Research, 2020, 48, 2676-2693.	14.5	30
43	The high protein expression of FOXO3, but not that of FOXO1, is associated with markers of good prognosis. Scientific Reports, 2020, 10, 6920.	3.3	5
44	ShallowHRD: detection of homologous recombination deficiency from shallow whole genome sequencing. Bioinformatics, 2020, 36, 3888-3889.	4.1	35
45	Biomarkers of cetuximab resistance in patients with head and neck squamous cell carcinoma. Cancer Biology and Medicine, 2020, 17, 208-217.	3.0	23
46	Response to mTOR and PI3K inhibitors in enzalutamide-resistant luminal androgen receptor triple-negative breast cancer patient-derived xenografts. Theranostics, 2020, 10, 1531-1543.	10.0	34
47	Differential gene expression in growth factors, epithelial mesenchymal transition and chemotaxis in the diffuse type compared with the intestinal type of gastric cancer. Oncology Letters, 2019, 18, 674-686.	1.8	18
48	P-cadherin-induced decorin secretion is required for collagen fiber alignment and directional collective cell migration. Journal of Cell Science, 2019, 132, .	2.0	14
49	Immune gene expression in head and neck squamous cell carcinoma patients. European Journal of Cancer, 2019, 121, 210-223.	2.8	45
50	Loss of the deglutamylase CCP5 perturbs multiple steps of spermatogenesis and leads to male infertility. Journal of Cell Science, 2019, 132, .	2.0	25
51	Clinical Development of Molecular Targeted Therapy in Head and Neck Squamous Cell Carcinoma. JNCI Cancer Spectrum, 2019, 3, pkz055.	2.9	34
52	PLEKHS1: A new molecular marker predicting risk of progression of non‑muscle‑invasive bladder cancer. Oncology Letters, 2019, 18, 3471-3480.	1.8	10
53	High Positive Correlations between ANRIL and p16-CDKN2A/p15-CDKN2B/p14-ARF Gene Cluster Overexpression in Multi-Tumor Types Suggest Deregulated Activation of an ANRIL–ARF Bidirectional Promoter. Non-coding RNA, 2019, 5, 44.	2.6	21
54	Inhibition of PI3K pathway increases immune infiltrate in muscle-invasive bladder cancer. OncoImmunology, 2019, 8, e1581556.	4.6	68

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55	A large collection of integrated genomically characterized patientâ€derived xenografts highlighting the heterogeneity of tripleâ€negative breast cancer. International Journal of Cancer, 2019, 145, 1902-1912.	5.1	37
56	Cortical branched actin determines cell cycle progression. Cell Research, 2019, 29, 432-445.	12.0	64
57	Interaction between IGF2â€PI3K axis and cancerâ€associatedâ€fibroblasts promotes anal squamous carcinogenesis. International Journal of Cancer, 2019, 145, 1852-1859.	5.1	13
58	Mechanistic Signatures of Human Papillomavirus Insertions in Anal Squamous Cell Carcinomas. Cancers, 2019, 11, 1846.	3.7	19
59	High Prevalence of a Hotspot of Noncoding Somatic Mutations in Intron 6 of <i>GPR126</i> in Bladder Cancer. Molecular Cancer Research, 2019, 17, 469-475.	3.4	18
60	Clinical Validity of HPV Circulating Tumor DNA in Advanced Anal Carcinoma: An Ancillary Study to the Epitopes-HPV02 Trial. Clinical Cancer Research, 2019, 25, 2109-2115.	7.0	65
61	Capecitabine Efficacy Is Correlated with TYMP and RB1 Expression in PDX Established from Triple-Negative Breast Cancers. Clinical Cancer Research, 2018, 24, 2605-2615.	7.0	45
62	Involvement of Aryl hydrocarbon receptor in myelination and in human nerve sheath tumorigenesis. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E1319-E1328.	7.1	27
63	Morphology and genomic hallmarks of breast tumours developed by ATM deleterious variant carriers. Breast Cancer Research, 2018, 20, 28.	5.0	35
64	Relevance of a molecular tumour board (MTB) for patients' enrolment in clinical trials: experience of the Institut Curie. ESMO Open, 2018, 3, e000339.	4.5	37
65	Revisited analysis of a SHIVA 01 trial cohort using functional mutational analyses successfully predicted treatment outcome. Molecular Oncology, 2018, 12, 594-601.	4.6	3
66	Comprehensive clinical and molecular analyses of neuroendocrine carcinomas of the breast. Modern Pathology, 2018, 31, 68-82.	5.5	58
67	Changes in chromatin state reveal ARNT2 at a node of a tumorigenic transcription factor signature driving glioblastoma cell aggressiveness. Acta Neuropathologica, 2018, 135, 267-283.	7.7	19
68	Tumor <i>PIK3CA</i> Genotype and Prognosis in Early-Stage Breast Cancer: A Pooled Analysis of Individual Patient Data. Journal of Clinical Oncology, 2018, 36, 981-990.	1.6	95
69	Distinct expression profiles and functions of Kindlins in breast cancer. Journal of Experimental and Clinical Cancer Research, 2018, 37, 281.	8.6	14
70	VOPP1 promotes breast tumorigenesis by interacting with the tumor suppressor WWOX. BMC Biology, 2018, 16, 109.	3.8	26
71	High rate of <i><scp>PlK</scp>3<scp>CA</scp></i> mutations but no <i><scp>TP</scp>53</i> mutations in lowâ€grade adenosquamous carcinoma of the breast. Histopathology, 2018, 73, 273-283.	2.9	33
72	Array comparative genomic hybridization identifies high level of PI3K/Akt/mTOR pathway alterations in anal cancer recurrences. Cancer Medicine, 2018, 7, 3213-3225.	2.8	13

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73	The trimeric coiledâ€coil <scp>HSBP</scp> 1 protein promotes <scp>WASH</scp> complex assembly at centrosomes. EMBO Journal, 2018, 37, .	7.8	22
74	Coronin 1C promotes triple-negative breast cancer invasiveness through regulation of MT1-MMP traffic and invadopodia function. Oncogene, 2018, 37, 6425-6441.	5.9	36
75	ETV4 transcription factor and MMP13 metalloprotease are interplaying actors of breast tumorigenesis. Breast Cancer Research, 2018, 20, 73.	5.0	56
76	Prognostic Impact of Residual HPV ctDNA Detection after Chemoradiotherapy for Anal Squamous Cell Carcinoma. Clinical Cancer Research, 2018, 24, 5767-5771.	7.0	68
77	MT1-MMP targeting to endolysosomes is mediated by flotillin upregulation. Journal of Cell Science, 2018, 131, .	2.0	29
78	Targeted next-generation sequencing identifies clinically relevant somatic mutations in a large cohort of inflammatory breast cancer. Breast Cancer Research, 2018, 20, 88.	5.0	53
79	Medullary Breast Carcinoma, a Triple-Negative Breast Cancer Associated with BCLG Overexpression. American Journal of Pathology, 2018, 188, 2378-2391.	3.8	12
80	The iron chelator deferasirox synergises with chemotherapy to treat tripleâ€negative breast cancers. Journal of Pathology, 2018, 246, 103-114.	4.5	47
81	High AHR expression in breast tumors correlates with expression of genes from several signaling pathways namely inflammation and endogenous tryptophan metabolism. PLoS ONE, 2018, 13, e0190619.	2.5	69
82	Exome sequencing reveals aberrant signalling pathways as hallmark of treatment-naive anal squamous cell carcinoma. Oncotarget, 2018, 9, 464-476.	1.8	23
83	Involvement of the FOXO6 transcriptional factor in breast carcinogenesis. Oncotarget, 2018, 9, 7464-7475.	1.8	12
84	PKD1 is a potential biomarker and therapeutic target in triple-negative breast cancer. Oncotarget, 2018, 9, 23208-23219.	1.8	14
85	Inhibition of mTOR downregulates expression of DNA repair proteins and is highly efficient against BRCA2-mutated breast cancer in combination to PARP inhibition. Oncotarget, 2018, 9, 29587-29600.	1.8	18
86	COX2/PTGS2 Expression Is Predictive of Response to Neoadjuvant Celecoxib in HER2-negative Breast Cancer Patients. Anticancer Research, 2018, 38, 1485-1490.	1.1	9
87	First French Pilot Quality Assessment of the EndoPredict Test for Early Luminal Breast Carcinoma. Anticancer Research, 2018, 38, 2909-2914.	1.1	3
88	Alterations in the balance of tubulin glycylation and glutamylation in photoreceptors leads to retinal degeneration. Journal of Cell Science, 2017, 130, 938-949.	2.0	57
89	The critical role of the <scp>ZNF217</scp> oncogene in promoting breast cancer metastasis to the bone. Journal of Pathology, 2017, 242, 73-89.	4.5	42
90	Baseline β-catenin, programmed death-ligand 1 expression and tumour-infiltrating lymphocytes predict response and poor prognosis in BRAF inhibitor-treated melanoma patients. European Journal of Cancer, 2017, 78, 70-81.	2.8	42

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91	Clinical value of R-spondins in triple-negative and metaplastic breast cancers. British Journal of Cancer, 2017, 116, 1595-1603.	6.4	31
92	Metabolic Response to Everolimus in Patient-Derived Triple-Negative Breast Cancer Xenografts. Journal of Proteome Research, 2017, 16, 1868-1879.	3.7	17
93	Correlation between messenger RNA expression and protein expression of immune checkpoint–associated molecules in bladder urothelial carcinoma: A retrospective study. Urologic Oncology: Seminars and Original Investigations, 2017, 35, 257-263.	1.6	29
94	Feasibility and clinical integration of molecular profiling for target identification in pediatric solid tumors. Pediatric Blood and Cancer, 2017, 64, e26365.	1.5	56
95	Protein biomarkers predictive for response to anti-EGFR treatment in RAS wild-type metastatic colorectal carcinoma. British Journal of Cancer, 2017, 117, 1819-1827.	6.4	15
96	mRNA Expression levels of genes involved in antitumor immunity: Identification of a 3-gene signature associated with prognosis of muscle-invasive bladder cancer. Oncolmmunology, 2017, 6, e1358330.	4.6	15
97	Tubulin glycylation controls primary cilia length. Journal of Cell Biology, 2017, 216, 2701-2713.	5.2	67
98	Inactivation of the Kinase Domain of CDK10 Prevents Tumor Growth in a Preclinical Model of Colorectal Cancer, and Is Accompanied by Downregulation of Bcl-2. Molecular Cancer Therapeutics, 2017, 16, 2292-2303.	4.1	14
99	Cytidine Deaminase Deficiency Reveals New Therapeutic Opportunities against Cancer. Clinical Cancer Research, 2017, 23, 2116-2126.	7.0	28
100	The SHIVA01 trial: what have we learned?. Pharmacogenomics, 2017, 18, 831-834.	1.3	4
101	Neuronal Cholesterol Accumulation Induced by Cyp46a1 Down-Regulation in Mouse Hippocampus Disrupts Brain Lipid Homeostasis. Frontiers in Molecular Neuroscience, 2017, 10, 211.	2.9	25
102	HPV circulating tumor DNA to monitor the efficacy of antiâ€PDâ€1 therapy in metastatic squamous cell carcinoma of the anal canal: A case report. International Journal of Cancer, 2017, 141, 1667-1670.	5.1	29
103	Combination of Carboplatin and Bevacizumab Is an Efficient Therapeutic Approach in Retinoblastoma Patient-Derived Xenografts. , 2016, 57, 4916.		10
104	Biopathological Significance of TLR9 Expression in Cancer Cells and Tumor Microenvironment Across Invasive Breast Carcinomas Subtypes. Cancer Microenvironment, 2016, 9, 107-118.	3.1	17
105	Vandetanib as a potential new treatment for estrogen receptorâ€negative breast cancers. International Journal of Cancer, 2016, 138, 2510-2521.	5.1	32
106	Prognostic value of a newly identified MALAT1 alternatively spliced transcript in breast cancer. British Journal of Cancer, 2016, 114, 1395-1404.	6.4	75
107	Expression of <i>ANRIL</i> –Polycomb Complexes– <i>CDKN2A/B/ARF</i> Genes in Breast Tumors: Identification of a Two-Gene (<i>EZH2/CBX7</i>) Signature with Independent Prognostic Value. Molecular Cancer Research, 2016, 14, 623-633.	3.4	84
108	Aryl hydrocarbon receptor–dependent enrichment of a megakaryocytic precursor with a high potential to produce proplatelets. Blood, 2016, 127, 2231-2240.	1.4	54

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109	Gain-of-function Prolactin Receptor Variants Are Not Associated With Breast Cancer and Multiple Fibroadenoma Risk. Journal of Clinical Endocrinology and Metabolism, 2016, 101, 4449-4460.	3.6	10
110	MicroRNA-based diagnostic tools for advanced fibrosis and cirrhosis in patients with chronic hepatitis B and C. Scientific Reports, 2016, 6, 34935.	3.3	41
111	Mutational analysis of anal cancers demonstrates frequent PIK3CA mutations associated with poor outcome after salvage abdominoperineal resection. British Journal of Cancer, 2016, 114, 1387-1394.	6.4	43
112	Treatment Algorithms Based on Tumor Molecular Profiling: The Essence of Precision Medicine Trials. Journal of the National Cancer Institute, 2016, 108, djv362.	6.3	71
113	CAG repeat size in Huntingtin alleles is associated with cancer prognosis. European Journal of Human Genetics, 2016, 24, 1310-1315.	2.8	19
114	Arpin downregulation in breast cancer is associated with poor prognosis. British Journal of Cancer, 2016, 114, 545-553.	6.4	25
115	Activation of IFN/STAT1 signalling predicts response to chemotherapy in oestrogen receptor-negative breast cancer. British Journal of Cancer, 2016, 114, 177-187.	6.4	67
116	p63/MT1-MMP axis is required for in situ to invasive transition in basal-like breast cancer. Oncogene, 2016, 35, 344-357.	5.9	76
117	Abstract CT041: Anti-proliferative response and predictive biomarkers to palbociclib in early breast cancer: The Preoperative Palbociclib (POP) randomized trial. , 2016, , .		10
118	PI3KCA mutation as an independent pronostic factor in anal squamous cell carcinoma treated by abdomino-perineal resection: Evidence from a retrospective cohort of 148 patients Journal of Clinical Oncology, 2016, 34, e15060-e15060.	1.6	1
119	Identification of new candidate therapeutic target genes in head and neck squamous cell carcinomas. Oncotarget, 2016, 7, 47418-47430.	1.8	13
120	Targeting mTOR pathway inhibits tumor growth in different molecular subtypes of triple-negative breast cancers. Oncotarget, 2016, 7, 48206-48219.	1.8	32
121	<i>ERBB2</i> mutations associated with solid variant of high-grade invasive lobular breast carcinomas. Oncotarget, 2016, 7, 73337-73346.	1.8	34
122	<i>MED12</i> mutations in breast phyllodes tumors: evidence of temporal tumoral heterogeneity and identification of associated critical signaling pathways. Oncotarget, 2016, 7, 84428-84438.	1.8	27
123	Combination of COX-2 expression and <i>PIK3CA</i> mutation as prognostic and predictive markers for celecoxib treatment in breast cancer. Oncotarget, 2016, 7, 85124-85141.	1.8	13
124	Everolimus affects vasculogenic mimicry in renal carcinoma resistant to sunitinib. Oncotarget, 2016, 7, 38467-38486.	1.8	31
125	mRNA expression levels and prognostic value of PD1/PDL1 and CTLA4 pathways genes in a large series of 155 bladder tumors Journal of Clinical Oncology, 2016, 34, 4523-4523.	1.6	0
126	Monitoring anti-PD-1 therapy efficacy by circulating tumor DNA: a prospective cohort Journal of Clinical Oncology, 2016, 34, 11535-11535.	1.6	0

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127	Biomarker-driven access to vemurafenib in BRAF-positive cancers: Second study of the French National AcSé Program Journal of Clinical Oncology, 2016, 34, TPS11620-TPS11620.	1.6	1
128	Abstract 1524: Mutations and gene copy number variations landscape of metastases of various cancer types from patients enrolled in the SHIVA trial. , 2016, , .		1
129	The GALNT9, BNC1 and CCDC8 genes are frequently epigenetically dysregulated in breast tumours that metastasise to the brain. Clinical Epigenetics, 2015, 7, 57.	4.1	75
130	IFI35, mir-99a and HCV Genotype to Predict Sustained Virological Response to Pegylated-Interferon Plus Ribavirin in Chronic Hepatitis C. PLoS ONE, 2015, 10, e0121395.	2.5	8
131	Long Noncoding RNAs as New Architects in Cancer Epigenetics, Prognostic Biomarkers, and Potential Therapeutic Targets. BioMed Research International, 2015, 2015, 1-14.	1.9	122
132	ATM has a major role in the double-strand break repair pathway dysregulation in sporadic breast carcinomas and is an independent prognostic marker at both mRNA and protein levels. British Journal of Cancer, 2015, 112, 1059-1066.	6.4	45
133	MiR-190b, the highest up-regulated miRNA in ERα-positive compared to ERα-negative breast tumors, a new biomarker in breast cancers?. BMC Cancer, 2015, 15, 499.	2.6	48
134	The Tumor-Suppressor WWOX and HDAC3 Inhibit the Transcriptional Activity of the \hat{l}^2 -Catenin Coactivator BCL9-2 in Breast Cancer Cells. Molecular Cancer Research, 2015, 13, 902-912.	3.4	18
135	Precision medicine: lessons learned from the SHIVA trial – Authors' reply. Lancet Oncology, The, 2015, 16, e581-e582.	10.7	13
136	Circulating tumor DNA as a nonâ€invasive substitute to metastasis biopsy for tumor genotyping and personalized medicine in a prospective trial across all tumor types. Molecular Oncology, 2015, 9, 783-790.	4.6	248
137	Alternative splicingâ€regulated protein of hepatitis B virus hacks the TNFâ€Î±â€stimulated signaling pathways and limits the extent of liver inflammation. FASEB Journal, 2015, 29, 1879-1889.	0.5	18
138	Evaluating Patient-Derived Colorectal Cancer Xenografts as Preclinical Models by Comparison with Patient Clinical Data. Cancer Research, 2015, 75, 1560-1566.	0.9	117
139	Cdk5 promotes DNA replication stress checkpoint activation through RPA-32 phosphorylation, and impacts on metastasis free survival in breast cancer patients. Cell Cycle, 2015, 14, 3066-3078.	2.6	24
140	MDA-MB-231 breast cancer cells overexpressing single VEGF isoforms display distinct colonisation characteristics. British Journal of Cancer, 2015, 113, 773-785.	6.4	12
141	Midkine Lacking Its Last 40 Amino Acids Acts on Endothelial and Neuroblastoma Tumor Cells and Inhibits Tumor Development. Molecular Cancer Therapeutics, 2015, 14, 213-224.	4.1	4
142	CYP46A1 inhibition, brain cholesterol accumulation and neurodegeneration pave the way for Alzheimer's disease. Brain, 2015, 138, 2383-2398.	7.6	163
143	Thrombin receptor PAR-1 activation on endothelial progenitor cells enhances chemotaxis-associated genes expression and leukocyte recruitment by a COX-2-dependent mechanism. Angiogenesis, 2015, 18, 347-359.	7.2	24
144	Intracerebral Gene Therapy Using AAVrh.10-hARSA Recombinant Vector to Treat Patients with Early-Onset Forms of Metachromatic Leukodystrophy: Preclinical Feasibility and Safety Assessments in Nonhuman Primates. Human Gene Therapy Clinical Development, 2015, 26, 113-124.	3.1	68

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145	<i>PIK3CA</i> Pathway Mutations Predictive of Poor Response Following Standard Radiochemotherapy ± Cetuximab in Cervical Cancer Patients. Clinical Cancer Research, 2015, 21, 2530-2537.	7.0	48
146	High-throughput pharmacogenetics identifies SLCO1A2 polymorphisms as candidates to elucidate the risk of febrile neutropenia in the breast cancer RAPP-01 trial. Breast Cancer Research and Treatment, 2015, 153, 383-389.	2.5	11
147	MAP3K8/TPL-2/COT is a potential predictive marker for MEK inhibitor treatment in high-grade serous ovarian carcinomas. Nature Communications, 2015, 6, 8583.	12.8	42
148	Unraveling the Role of Huntingtin in Breast Cancer Metastasis. Journal of the National Cancer Institute, 2015, 107, djv208.	6.3	32
149	Cholesterol 24-hydroxylase defect is implicated in memory impairments associated with Alzheimer-like Tau pathology. Human Molecular Genetics, 2015, 24, 5965-5976.	2.9	96
150	Molecularly targeted therapy based on tumour molecular profiling versus conventional therapy for advanced cancer (SHIVA): a multicentre, open-label, proof-of-concept, randomised, controlled phase 2 trial. Lancet Oncology, The, 2015, 16, 1324-1334.	10.7	897
151	SWI/SNF Chromatin Remodeling and Human Malignancies. Annual Review of Pathology: Mechanisms of Disease, 2015, 10, 145-171.	22.4	242
152	Abstract 1687: Vandetanib as a potential new treatment for ER negative breast cancers. Cancer Research, 2015, 75, 1687-1687.	0.9	4
153	Tumor PIK3CA genotype and prognosis: A pooled analysis of 4,241 patients (pts) with early-stage breast cancer (BC) Journal of Clinical Oncology, 2015, 33, 516-516.	1.6	5
154	IL-1β produced by aggressive breast cancer cells is one of the factors that dictate their interactions with mesenchymal stem cells through chemokine production. Oncotarget, 2015, 6, 29034-29047.	1.8	56
155	Abstract P2-05-06: Identification of specific gene signatures and alternative splice variants using exon array in Inflammatory breast Cancer. , 2015, , .		0
156	Assessment of functional impact of germline BRCA1/2 variants located in noncoding regions in families with breast-ovarian cancer predisposition Journal of Clinical Oncology, 2015, 33, 1542-1542.	1.6	0
157	Abstract 2400: Circulating tumor DNA as a non-invasive substitute to metastasis biopsy for targeted sequencing in a prospective randomized trial for personalized treatment in all tumor type: The SHIVA study. , 2015, , .		0
158	Design and statistical principles of the SHIVA trial. Chinese Clinical Oncology, 2015, 4, 32.	1.2	1
159	Predictive Gene Signature of Response to the Anti-TweakR mAb PDL192 in Patient-Derived Breast Cancer Xenografts. PLoS ONE, 2014, 9, e104227.	2.5	10
160	Cooperation between human fibrocytes and endothelial colony-forming cells increases angiogenesis via the CXCR4 pathway. Thrombosis and Haemostasis, 2014, 112, 1002-1013.	3.4	30
161	Concomitant Notch activation and p53 deletion trigger epithelial-to-mesenchymal transition and metastasis in mouse gut. Nature Communications, 2014, 5, 5005.	12.8	114
162	Reduction of MicroRNA 122 Expression in <i>IFNL3</i> CT/TT Carriers and during Progression of Fibrosis in Patients with Chronic Hepatitis C. Journal of Virology, 2014, 88, 6394-6402.	3.4	29

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