Sarah-Jane Dawson

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Analysis of Circulating Tumor DNA to Monitor Metastatic Breast Cancer. New England Journal of Medicine, 2013, 368, 1199-1209.	13.9	1,884
2	Non-invasive analysis of acquired resistance to cancer therapy by sequencing of plasma DNA. Nature, 2013, 497, 108-112.	13.7	1,443
3	The somatic mutation profiles of 2,433 breast cancers refine their genomic and transcriptomic landscapes. Nature Communications, 2016, 7, 11479.	5.8	1,221
4	Noninvasive Identification and Monitoring of Cancer Mutations by Targeted Deep Sequencing of Plasma DNA. Science Translational Medicine, 2012, 4, 136ra68.	5.8	1,086
5	Clinical validity of circulating tumour cells in patients with metastatic breast cancer: a pooled analysis of individual patient data. Lancet Oncology, The, 2014, 15, 406-414.	5.1	703
6	BET inhibitor resistance emerges from leukaemia stem cells. Nature, 2015, 525, 538-542.	13.7	441
7	Multifocal clonal evolution characterized using circulating tumour DNA in a case of metastatic breast cancer. Nature Communications, 2015, 6, 8760.	5.8	409
8	An Evolutionarily Conserved Function of Polycomb Silences the MHC Class I Antigen Presentation Pathway and Enables Immune Evasion in Cancer. Cancer Cell, 2019, 36, 385-401.e8.	7.7	359
9	Ibrutinib plus Venetoclax for the Treatment of Mantle-Cell Lymphoma. New England Journal of Medicine, 2018, 378, 1211-1223.	13.9	343
10	Non-genetic mechanisms of therapeutic resistance in cancer. Nature Reviews Cancer, 2020, 20, 743-756.	12.8	290
11	Selective targeting of BD1 and BD2 of the BET proteins in cancer and immunoinflammation. Science, 2020, 368, 387-394.	6.0	274
12	A new genome-driven integrated classification of breast cancer and its implications. EMBO Journal, 2013, 32, 617-628.	3.5	267
13	The clinical use of circulating tumor cells (CTCs) enumeration for staging of metastatic breast cancer (MBC): International expert consensus paper. Critical Reviews in Oncology/Hematology, 2019, 134, 39-45.	2.0	200
14	Multi-omic machine learning predictor of breast cancer therapy response. Nature, 2022, 601, 623-629.	13.7	187
15	Reversion of <i>BRCA1/2</i> Germline Mutations Detected in Circulating Tumor DNA From Patients With High-Grade Serous Ovarian Cancer. Journal of Clinical Oncology, 2017, 35, 1274-1280.	0.8	157
16	Dynamic molecular monitoring reveals that SWI–SNF mutations mediate resistance to ibrutinib plus venetoclax in mantle cell lymphoma. Nature Medicine, 2019, 25, 119-129.	15.2	147
17	Targeting enhancer switching overcomes non-genetic drug resistance in acute myeloid leukaemia. Nature Communications, 2019, 10, 2723.	5.8	126
18	Click chemistry enables preclinical evaluation of targeted epigenetic therapies. Science, 2017, 356, 1397-1401.	6.0	120

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19	The Subclonal Architecture of Metastatic Breast Cancer: Results from a Prospective Community-Based Rapid Autopsy Program "CASCADE― PLoS Medicine, 2016, 13, e1002204.	3.9	119
20	HBO1 is required for the maintenance of leukaemia stem cells. Nature, 2020, 577, 266-270.	13.7	105
21	A Phase Ib Dose-Escalation and Expansion Study of the BCL2 Inhibitor Venetoclax Combined with Tamoxifen in ER and BCL2–Positive Metastatic Breast Cancer. Cancer Discovery, 2019, 9, 354-369.	7.7	104
22	Functional interdependence of BRD4 and DOT1L in MLL leukemia. Nature Structural and Molecular Biology, 2016, 23, 673-681.	3.6	92
23	The value of cellâ€free DNA for molecular pathology. Journal of Pathology, 2018, 244, 616-627.	2.1	91
24	Effects of Collection and Processing Procedures on Plasma Circulating Cell-Free DNA from Cancer Patients. Journal of Molecular Diagnostics, 2018, 20, 883-892.	1.2	81
25	Non-genetic determinants of malignant clonal fitness at single-cell resolution. Nature, 2022, 601, 125-131.	13.7	71
26	Circulating tumour DNA reflects treatment response and clonal evolution in chronic lymphocytic leukaemia. Nature Communications, 2017, 8, 14756.	5.8	70
27	A community-based model of rapid autopsy in end-stage cancer patients. Nature Biotechnology, 2016, 34, 1010-1014.	9.4	66
28	HIV is associated with an increased risk of age-related clonal hematopoiesis among older adults. Nature Medicine, 2021, 27, 1006-1011.	15.2	62
29	Molecular disease monitoring using circulating tumor DNA in myelodysplastic syndromes. Blood, 2017, 129, 1685-1690.	0.6	53
30	Circulating Tumor DNA Analysis and Functional Imaging Provide Complementary Approaches for Comprehensive Disease Monitoring in Metastatic Melanoma. JCO Precision Oncology, 2017, 1, 1-14.	1.5	51
31	Sustained clinical responses to tyrosine kinase inhibitor sunitinib in thyroid carcinoma. Anti-Cancer Drugs, 2008, 19, 547-552.	0.7	48
32	Evolution of late-stage metastatic melanoma is dominated by aneuploidy and whole genome doubling. Nature Communications, 2021, 12, 1434.	5.8	46
33	Coding and noncoding drivers of mantle cell lymphoma identified through exome and genome sequencing. Blood, 2020, 136, 572-584.	0.6	44
34	Towards Routine Implementation of Liquid Biopsies in Cancer Management: It Is Always Too Early, until Suddenly It Is Too Late. Diagnostics, 2021, 11, 103.	1.3	33
35	Three Year Update of the Phase II ABT-199 (Venetoclax) and Ibrutinib in Mantle Cell Lymphoma (AIM) Study. Blood, 2019, 134, 756-756.	0.6	24
36	Detection of cell-free microbial DNA using a contaminant-controlled analysis framework. Genome Biology, 2021, 22, 187.	3.8	22

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37	Cancer Risk Management Practices of Noncarriers Within <i>BRCA1/2</i> Mutation–Positive Families in the Kathleen Cuningham Foundation Consortium for Research Into Familial Breast Cancer. Journal of Clinical Oncology, 2008, 26, 225-232.	0.8	19
38	Circulating tumour DNA in metastatic breast cancer to guide clinical trial enrolment and precision oncology: AAcohort study. PLoS Medicine, 2020, 17, e1003363.	3.9	18
39	Pharmacologic Reduction of Mitochondrial Iron Triggers a Noncanonical BAX/BAK-Dependent Cell Death. Cancer Discovery, 2022, 12, 774-791.	7.7	18
40	Alpelisib Monotherapy for PI3K-Altered, Pretreated Advanced Breast Cancer: A Phase II Study. Cancer Discovery, 2022, 12, 2058-2073.	7.7	16
41	Liquid biopsies for residual disease and recurrence. Med, 2021, 2, 1292-1313.	2.2	15
42	Wet or Dry? Do Liquid Biopsy Techniques Compete with or Complement PET for Disease Monitoring in Oncology?. Journal of Nuclear Medicine, 2017, 58, 869-870.	2.8	8
43	Characterizing the Cancer Genome in Blood. Cold Spring Harbor Perspectives in Medicine, 2019, 9, a026880.	2.9	7
44	Combining liquid biopsies and PET-CT for early cancer detection. Nature Medicine, 2020, 26, 1010-1011.	15.2	7
45	Hypocalcemia associated with bone metastases in a patient with salivary-gland carcinoma. Nature Clinical Practice Oncology, 2006, 3, 104-107.	4.3	5
46	Blood Worth Bottling: Circulating Tumor DNA as a Cancer Biomarker. Cancer Research, 2016, 76, 5590-5591.	0.4	5
47	Modeling the Prognostic Impact of Circulating Tumor Cells Enumeration in Metastatic Breast Cancer for Clinical Trial Design Simulation. Oncologist, 2022, 27, e561-e570.	1.9	5
48	Large B-cell lymphoma: is the future written in the blood?. Lancet Oncology, The, 2015, 16, 481-483.	5.1	3
49	Plasma and tumor genomic correlates of response to BYL719 in PI3KCA mutated metastatic ER-positive breast cancer (ER+/HER2- BC) Journal of Clinical Oncology, 2018, 36, 1055-1055.	0.8	3
50	Clinical validation and implementation of droplet digital PCR for the detection of BRAF mutations from cell-free DNA. Pathology, 2022, 54, 772-778.	0.3	2
51	Custom workflows to improve joint variant calling from multiple related tumour samples: FreeBayesSomatic and Strelka2Pass. Bioinformatics, 2021, , .	1.8	Ο