

# Alan Ashworth

## List of Publications by Year in descending order

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Version: 2024-02-01

61  
papers

31,838  
citations

57758

44  
h-index

123424

61  
g-index

67  
all docs

67  
docs citations

67  
times ranked

35234  
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeting the DNA repair defect in BRCA mutant cells as a therapeutic strategy. Nature, 2005, 434, 917-921.	27.8	5,595
2	A SARS-CoV-2 protein interaction map reveals targets for drug repurposing. Nature, 2020, 583, 459-468.	27.8	3,542
3	Inhibition of Poly(ADP-Ribose) Polymerase in Tumors from BRCA Mutation Carriers. New England Journal of Medicine, 2009, 361, 123-134.	27.0	3,312
4	Identification of the breast cancer susceptibility gene BRCA2. Nature, 1995, 378, 789-792.	27.8	3,230
5	PARP inhibitors: Synthetic lethality in the clinic. Science, 2017, 355, 1152-1158.	12.6	1,826
6	DNA-Repair Defects and Olaparib in Metastatic Prostate Cancer. New England Journal of Medicine, 2015, 373, 1697-1708.	27.0	1,796
7	Hallmarks of 'BRCAness' in sporadic cancers. Nature Reviews Cancer, 2004, 4, 814-819.	28.4	1,477
8	Deficiency in the Repair of DNA Damage by Homologous Recombination and Sensitivity to Poly(ADP-Ribose) Polymerase Inhibition. Cancer Research, 2006, 66, 8109-8115.	0.9	1,172
9	BRCAness revisited. Nature Reviews Cancer, 2016, 16, 110-120.	28.4	976
10	Resistance to therapy caused by intragenic deletion in BRCA2. Nature, 2008, 451, 1111-1115.	27.8	894
11	Poly(ADP)-Ribose Polymerase Inhibition: Frequent Durable Responses in BRCA Carrier Ovarian Cancer Correlating With Platinum-Free Interval. Journal of Clinical Oncology, 2010, 28, 2512-2519.	1.6	877
12	Carboplatin in BRCA1/2-mutated and triple-negative breast cancer BRCAness subgroups: the TNT Trial. Nature Medicine, 2018, 24, 628-637.	30.7	649
13	Genomic Hallmarks and Structural Variation in Metastatic Prostate Cancer. Cell, 2018, 174, 758-769.e9.	28.9	459
14	BMN 673, a Novel and Highly Potent PARP1/2 Inhibitor for the Treatment of Human Cancers with DNA Repair Deficiency. Clinical Cancer Research, 2013, 19, 5003-5015.	7.0	416
15	Genetic Interactions in Cancer Progression and Treatment. Cell, 2011, 145, 30-38.	28.9	380
16	Genome-wide CRISPR Screens in Primary Human T Cells Reveal Key Regulators of Immune Function. Cell, 2018, 175, 1958-1971.e15.	28.9	378
17	Mechanisms of resistance to therapies targeting BRCA-mutant cancers. Nature Medicine, 2013, 19, 1381-1388.	30.7	371
18	Genome-wide and high-density CRISPR-Cas9 screens identify point mutations in PARP1 causing PARP inhibitor resistance. Nature Communications, 2018, 9, 1849.	12.8	310

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19	Synthetic lethal therapies for cancer: what's next after PARP inhibitors?. <i>Nature Reviews Clinical Oncology</i> , 2018, 15, 564-576.	27.6	303
20	Synthetic lethality as an engine for cancer drug target discovery. <i>Nature Reviews Drug Discovery</i> , 2020, 19, 23-38.	46.4	295
21	A Marker of Homologous Recombination Predicts Pathologic Complete Response to Neoadjuvant Chemotherapy in Primary Breast Cancer. <i>Clinical Cancer Research</i> , 2010, 16, 6159-6168.	7.0	287
22	Secondary mutations in <i>BRCA2</i> associated with clinical resistance to a <i>PARP</i> inhibitor. <i>Journal of Pathology</i> , 2013, 229, 422-429.	4.5	287
23	ATR inhibitors as a synthetic lethal therapy for tumours deficient in <i>ARID1A</i> . <i>Nature Communications</i> , 2016, 7, 13837.	12.8	272
24	PARP inhibition enhances tumor cell's intrinsic immunity in <i>ERCC1</i> -deficient non-small cell lung cancer. <i>Journal of Clinical Investigation</i> , 2019, 129, 1211-1228.	8.2	222
25	PARP inhibitor combination therapy. <i>Critical Reviews in Oncology/Hematology</i> , 2016, 108, 73-85.	4.4	175
26	Transcription-Associated Cyclin-Dependent Kinases as Targets and Biomarkers for Cancer Therapy. <i>Cancer Discovery</i> , 2020, 10, 351-370.	9.4	162
27	ADP-ribosyltransferases, an update on function and nomenclature. <i>FEBS Journal</i> , 2022, 289, 7399-7410.	4.7	150
28	A Genetic Screen Using the PiggyBac Transposon in Haploid Cells Identifies <i>Parp1</i> as a Mediator of Olaparib Toxicity. <i>PLoS ONE</i> , 2013, 8, e61520.	2.5	147
29	An expanded universe of cancer targets. <i>Cell</i> , 2021, 184, 1142-1155.	28.9	135
30	<i>MKP5</i> , a new member of the MAP kinase phosphatase family, which selectively dephosphorylates stress-activated kinases. <i>Oncogene</i> , 1999, 18, 6981-6988.	5.9	132
31	A Short Pseudoautosomal Region in Laboratory Mice. <i>Genome Research</i> , 2001, 11, 1826-1832.	5.5	120
32	The Cancer Cell Map Initiative: Defining the Hallmark Networks of Cancer. <i>Molecular Cell</i> , 2015, 58, 690-698.	9.7	117
33	Fragment binding to the Nsp3 macrodomain of SARS-CoV-2 identified through crystallographic screening and computational docking. <i>Science Advances</i> , 2021, 7, .	10.3	100
34	Large-Scale Profiling of Kinase Dependencies in Cancer Cell Lines. <i>Cell Reports</i> , 2016, 14, 2490-2501.	6.4	97
35	Evolutionary rate of a gene affected by chromosomal position. <i>Current Biology</i> , 1999, 9, 987-S3.	3.9	94
36	Targeting DNA Damage Response and Replication Stress in Pancreatic Cancer. <i>Gastroenterology</i> , 2021, 160, 362-377.e13.	1.3	90

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37	Elevated APOBEC3B expression drives a kataegic-like mutation signature and replication stress-related therapeutic vulnerabilities in p53-defective cells. <i>British Journal of Cancer</i> , 2017, 117, 113-123.	6.4	84
38	Commonly Occurring Cell Subsets in High-Grade Serous Ovarian Tumors Identified by Single-Cell Mass Cytometry. <i>Cell Reports</i> , 2018, 22, 1875-1888.	6.4	83
39	HNF4A and GATA6 Loss Reveals Therapeutically Actionable Subtypes in Pancreatic Cancer. <i>Cell Reports</i> , 2020, 31, 107625.	6.4	78
40	Complementation of byr1 in fission yeast by mammalian MAP kinase kinase requires coexpression of Raf kinase. <i>Nature</i> , 1993, 364, 349-352.	27.8	76
41	Synthetic Lethal Targeting of <i>ARID1A</i> -Mutant Ovarian Clear Cell Tumors with Dasatinib. <i>Molecular Cancer Therapeutics</i> , 2016, 15, 1472-1484.	4.1	73
42	Modeling Therapy Resistance in <i>BRCA1/2</i> -Mutant Cancers. <i>Molecular Cancer Therapeutics</i> , 2017, 16, 2022-2034.	4.1	66
43	A protein interaction landscape of breast cancer. <i>Science</i> , 2021, 374, eabf3066.	12.6	66
44	CDK1 Is a Synthetic Lethal Target for KRAS Mutant Tumours. <i>PLoS ONE</i> , 2016, 11, e0149099.	2.5	60
45	Functional Genetic Screen Identifies Increased Sensitivity to WEE1 Inhibition in Cells with Defects in Fanconi Anemia and HR Pathways. <i>Molecular Cancer Therapeutics</i> , 2015, 14, 865-876.	4.1	52
46	ATR Is a Therapeutic Target in Synovial Sarcoma. <i>Cancer Research</i> , 2017, 77, 7014-7026.	0.9	43
47	A novel tankyrase inhibitor, MSC2504877, enhances the effects of clinical CDK4/6 inhibitors. <i>Scientific Reports</i> , 2019, 9, 201.	3.3	38
48	Assessing the Significance of <i>BRCA1</i> and <i>BRCA2</i> Mutations in Pancreatic Cancer. <i>Journal of Clinical Oncology</i> , 2015, 33, 3080-3081.	1.6	31
49	Design and Synthesis of Poly(ADP-ribose) Polymerase Inhibitors: Impact of Adenosine Pocket-Binding Motif Appendage to the 3-Oxo-2,3-dihydrobenzofuran-7-carboxamide on Potency and Selectivity. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 5330-5357.	6.4	26
50	The cylindromatosis gene product, CYLD, interacts with MIB2 to regulate Notch signalling. <i>Oncotarget</i> , 2014, 5, 12126-12140.	1.8	26
51	An in vivo model of intratumoural aromatase using aromatase-transfected MCF7 human breast cancer cells. <i>International Journal of Cancer</i> , 1995, 62, 297-302.	5.1	24
52	The mechanisms of catalysis and ligand binding for the SARS-CoV-2 NSP3 macrodomain from neutron and x-ray diffraction at room temperature. <i>Science Advances</i> , 2022, 8, .	10.3	24
53	Coupling bimolecular PARylation biosensors with genetic screens to identify PARylation targets. <i>Nature Communications</i> , 2018, 9, 2016.	12.8	22
54	DNA repair deficiency sensitizes lung cancer cells to NAD <sup>+</sup> biosynthesis blockade. <i>Journal of Clinical Investigation</i> , 2018, 128, 1671-1687.	8.2	19

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55	Complementary genetic screens identify the E3 ubiquitin ligase CBL, as a modifier of PARP inhibitor sensitivity. <i>Oncotarget</i> , 2015, 6, 10746-10758.	1.8	16
56	Genome-wide barcoded transposon screen for cancer drug sensitivity in haploid mouse embryonic stem cells. <i>Scientific Data</i> , 2017, 4, 170020.	5.3	14
57	Chemosensitivity profiling of osteosarcoma tumour cell lines identifies a model of BRCAness. <i>Scientific Reports</i> , 2018, 8, 10614.	3.3	13
58	Oncogenic KRAS sensitizes premalignant, but not malignant cells, to Noxa-dependent apoptosis through the activation of the MEK/ERK pathway. <i>Oncotarget</i> , 2015, 6, 10994-11008.	1.8	13
59	A Very Long-Acting PARP Inhibitor Suppresses Cancer Cell Growth in DNA Repair-Deficient Tumor Models. <i>Cancer Research</i> , 2021, 81, 1076-1086.	0.9	10
60	A Whole-Genome CRISPR Screen Identifies AHR Loss as a Mechanism of Resistance to a PARP7 Inhibitor. <i>Molecular Cancer Therapeutics</i> , 2022, 21, 1076-1089.	4.1	8
61	Hypoxia Is a Dominant Remodeler of the Effector T Cell Surface Proteome Relative to Activation and Regulatory T Cell Suppression. <i>Molecular and Cellular Proteomics</i> , 2022, 21, 100217.	3.8	5