

Thomas P Howard

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4041917/publications.pdf>

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13
papers

1,046
citations

933447

10
h-index

1125743

13
g-index

23
all docs

23
docs citations

23
times ranked

2808
citing authors

#	ARTICLE	IF	CITATIONS
1	A first-generation pediatric cancer dependency map. <i>Nature Genetics</i> , 2021, 53, 529-538.	21.4	76
2	Haplotype-resolved germline and somatic alterations in renal medullary carcinomas. <i>Genome Medicine</i> , 2021, 13, 114.	8.2	5
3	Clinical outcomes for pediatric patients receiving radiotherapy for solid tumor central nervous system metastases. <i>Pediatric Blood and Cancer</i> , 2021, 68, e29331.	1.5	1
4	Rhabdoid Tumors Are Sensitive to the Protein-Translation Inhibitor Homoharringtonine. <i>Clinical Cancer Research</i> , 2020, 26, 4995-5006.	7.0	14
5	Partitioning of Chemotherapeutics into Nuclear Condensates—Opening the Door to New Approaches for Drug Development. <i>Molecular Cell</i> , 2020, 79, 544-545.	9.7	7
6	BRD9 defines a SWI/SNF sub-complex and constitutes a specific vulnerability in malignant rhabdoid tumors. <i>Nature Communications</i> , 2019, 10, 1881.	12.8	117
7	MDM2 and MDM4 Are Therapeutic Vulnerabilities in Malignant Rhabdoid Tumors. <i>Cancer Research</i> , 2019, 79, 2404-2414.	0.9	43
8	Renal medullary carcinomas depend upon SMARCB1 loss and are sensitive to proteasome inhibition. <i>ELife</i> , 2019, 8, .	6.0	32
9	Mutational processes shape the landscape of TP53 mutations in human cancer. <i>Nature Genetics</i> , 2018, 50, 1381-1387.	21.4	334
10	Functional Genomic Characterization of Cancer Genomes. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2016, 81, 237-246.	1.1	17
11	Control of sexuality by the <i>sk1</i> -encoded UDP-glycosyltransferase of maize. <i>Science Advances</i> , 2016, 2, e1600991.	10.3	37
12	SWI/SNF-mutant cancers depend on catalytic and non-catalytic activity of EZH2. <i>Nature Medicine</i> , 2015, 21, 1491-1496.	30.7	334
13	Identification of the Maize Gravitropism Gene <i>lazy plant1</i> by a Transposon-Tagging Genome Resequencing Strategy. <i>PLoS ONE</i> , 2014, 9, e87053.	2.5	25