Jason R Dobson

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3499594/publications.pdf

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21 papers

4,231 citations

20 h-index 713466 21 g-index

21 all docs

21 docs citations

times ranked

21

9878 citing authors

#	Article	IF	CITATIONS
1	Phase 1 study of single-agent WNT974, a first-in-class Porcupine inhibitor, in patients with advanced solid tumours. British Journal of Cancer, 2021, 125, 28-37.	6.4	62
2	A Phase I Study of the CDK4/6 Inhibitor Ribociclib (LEEO11) in Pediatric Patients with Malignant Rhabdoid Tumors, Neuroblastoma, and Other Solid Tumors. Clinical Cancer Research, 2017, 23, 2433-2441.	7.0	134
3	Genomic Analysis of Nasopharyngeal Carcinoma Reveals TME-Based Subtypes. Molecular Cancer Research, 2017, 15, 1722-1732.	3.4	119
4	Identifying Nuclear Matrixâ€Attached DNA Across the Genome. Journal of Cellular Physiology, 2017, 232, 1295-1305.	4.1	19
5	Runx1 stabilizes the mammary epithelial cell phenotype and prevents epithelial to mesenchymal transition. Oncotarget, 2017, 8, 17610-17627.	1.8	53
6	A Phase I Study of the Cyclin-Dependent Kinase 4/6 Inhibitor Ribociclib (LEE011) in Patients with Advanced Solid Tumors and Lymphomas. Clinical Cancer Research, 2016, 22, 5696-5705.	7.0	245
7	Allosteric inhibition of SHP2 phosphatase inhibits cancers driven by receptor tyrosine kinases. Nature, 2016, 535, 148-152.	27.8	674
8	Expansion of GA Dinucleotide Repeats Increases the Density of CLAMP Binding Sites on the X-Chromosome to Promote Drosophila Dosage Compensation. PLoS Genetics, 2016, 12, e1006120.	3.5	48
9	The BRG1 chromatin remodeling enzyme links cancer cell metabolism and proliferation. Oncotarget, 2016, 7, 38270-38281.	1.8	51
10	Expression of the ILâ€11 Gene in Metastatic Cells Is Supported by Runx2â€5mad and Runx2â€cJun Complexes Induced by TGFβ1. Journal of Cellular Biochemistry, 2015, 116, 2098-2108.	2.6	21
11	The SWI/SNF ATPases Are Required for Triple Negative Breast Cancer Cell Proliferation. Journal of Cellular Physiology, 2015, 230, 2683-2694.	4.1	58
12	Pan-cancer network analysis identifies combinations of rare somatic mutations across pathways and protein complexes. Nature Genetics, 2015, 47, 106-114.	21.4	830
13	The bone-specific Runx2-P1 promoter displays conserved three-dimensional chromatin structure with the syntenic Supt3h promoter. Nucleic Acids Research, 2014, 42, 10360-10372.	14.5	28
14	Genomic occupancy of Runx2 with global expression profiling identifies a novel dimension to control of osteoblastogenesis. Genome Biology, 2014, 15, R52.	9.6	122
15	Identifying driver mutations in sequenced cancer genomes: computational approaches to enable precision medicine. Genome Medicine, 2014, 6, 5.	8.2	186
16	Multiplatform Analysis of 12 Cancer Types Reveals Molecular Classification within and across Tissues of Origin. Cell, 2014, 158, 929-944.	28.9	1,242
17	hsa-mir-30c promotes the invasive phenotype of metastatic breast cancer cells by targeting NOV/CCN3. Cancer Cell International, 2014, 14, 73.	4.1	46
18	A Runx2-HDAC1 co-repressor complex regulates rRNA gene expression by modulating UBF acetylation. Journal of Cell Science, 2012, 125, 2732-9.	2.0	36

#	Article	IF	CITATIONS
19	Transcriptional corepressor TLE1 functions with Runx2 in epigenetic repression of ribosomal RNA genes. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 4165-4169.	7.1	41
20	Cancer-related ectopic expression of the bone-related transcription factor RUNX2 in non-osseous metastatic tumor cells is linked to cell proliferation and motility. Breast Cancer Research, 2010, 12, R89.	5.0	56
21	Runx2 Transcriptional Activation of Indian Hedgehog and a Downstream Bone Metastatic Pathway in Breast Cancer Cells. Cancer Research, 2008, 68, 7795-7802.	0.9	160