Ian J Davis

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3138561/publications.pdf

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31	1,514	20	29
papers	citations	h-index	g-index
31	31	31	3430
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	PBRM1 Inactivation Promotes Upregulation of Human Endogenous Retroviruses in a HIF-Dependent Manner. Cancer Immunology Research, 2022, 10, 285-290.	3.4	13
2	Targeting disialoganglioside GD2 with chimeric antigen receptor-redirected T cells in lung cancer. , 2022, 10, e003897.		27
3	HDAC inhibition results in widespread alteration of the histone acetylation landscape and BRD4 targeting to gene bodies. Cell Reports, 2021, 34, 108638.	6.4	60
4	Genome-wide cancer-specific chromatin accessibility patterns derived from archival processed xenograft tumors. Genome Research, 2021, 31, 2327-2339.	5.5	3
5	An optogenetic switch for the Set2 methyltransferase provides evidence for transcription-dependent and -independent dynamics of H3K36 methylation. Genome Research, 2020, 30, 1605-1617.	5.5	10
6	Accelerated aging among childhood, adolescent, and young adult cancer survivors is evidenced by increased expression of p16 ^{INK4a} and frailty. Cancer, 2020, 126, 4975-4983.	4.1	47
7	Discrete Adaptive Responses to MEK Inhibitor in Subpopulations of Triple-Negative Breast Cancer. Molecular Cancer Research, 2020, 18, 1685-1698.	3.4	3
8	Emerging novel agents for patients with advanced Ewing sarcoma: a report from the Children's Oncology Group (COG) New Agents for Ewing Sarcoma Task Force. F1000Research, 2019, 8, 493.	1.6	57
9	EWSR1-FLI1 Activation of the Cancer/Testis Antigen FATE1 Promotes Ewing Sarcoma Survival. Molecular and Cellular Biology, 2019, 39, .	2.3	15
10	Chromatin Accessibility as a Strategy to Detect Changes Associated With Development, Disease, and Exposure and Susceptibility to Chemical Toxins., 2019,, 85-103.		3
11	Cavitation Enhancement Increases the Efficiency and Consistency of Chromatin Fragmentation from Fixed Cells for Downstream Quantitative Applications. Biochemistry, 2018, 57, 2756-2761.	2.5	11
12	Casein Kinase II Phosphorylation of Spt6 Enforces Transcriptional Fidelity by Maintaining Spn1-Spt6 Interaction. Cell Reports, 2018, 25, 3476-3489.e5.	6.4	20
13	Spt6 Association with RNA Polymerase II Directs mRNA Turnover During Transcription. Molecular Cell, 2018, 70, 1054-1066.e4.	9.7	38
14	H3K36 Methylation Regulates Nutrient Stress Response in Saccharomyces cerevisiae by Enforcing Transcriptional Fidelity. Cell Reports, 2017, 19, 2371-2382.	6.4	54
15	Radiation Sensitivity in a Preclinical Mouse Model of Medulloblastoma Relies on the Function of the Intrinsic Apoptotic Pathway. Cancer Research, 2016, 76, 3211-3223.	0.9	25
16	Dual Chromatin and Cytoskeletal Remodeling by SETD2. Cell, 2016, 166, 950-962.	28.9	204
17	Structure/Function Analysis of Recurrent Mutations in SETD2 Protein Reveals a Critical and Conserved Role for a SET Domain Residue in Maintaining Protein Stability and Histone H3 Lys-36 Trimethylation. Journal of Biological Chemistry, 2016, 291, 21283-21295.	3.4	64
18	Widespread Chromatin Accessibility at Repetitive Elements Links Stem Cells with Human Cancer. Cell Reports, 2016, 17, 1607-1620.	6.4	32

#	Article	IF	CITATIONS
19	Post-transplant lymphoproliferative disorder of the pediatric airway: Presentation and management. International Journal of Pediatric Otorhinolaryngology, 2016, 86, 218-223.	1.0	9
20	Subtumoral analysis of PRINT nanoparticle distribution reveals targeting variation based on cellular and particle properties. Nanomedicine: Nanotechnology, Biology, and Medicine, 2016, 12, 1053-1062.	3.3	27
21	The Cardiac TBX5 Interactome Reveals a Chromatin Remodeling Network Essential for Cardiac Septation. Developmental Cell, 2016, 36, 262-275.	7.0	71
22	High-throughput small molecule screen identifies inhibitors of aberrant chromatin accessibility. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 3018-3023.	7.1	26
23	A Role for Widely Interspaced Zinc Finger (WIZ) in Retention of the G9a Methyltransferase on Chromatin. Journal of Biological Chemistry, 2015, 290, 26088-26102.	3.4	29
24	Linking germline and somatic variation in Ewing sarcoma. Nature Genetics, 2015, 47, 964-965.	21.4	4
25	Variation in chromatin accessibility in human kidney cancer links H3K36 methyltransferase loss with widespread RNA processing defects. Genome Research, 2014, 24, 241-250.	5.5	160
26	LKB1 loss in melanoma disrupts directional migration toward extracellular matrix cues. Journal of Cell Biology, 2014, 207, 299-315.	5.2	41
27	PTEN Deficiency Mediates a Reciprocal Response to IGFI and mTOR Inhibition. Molecular Cancer Research, 2014, 12, 1610-1620.	3.4	25
28	A Detailed Protocol for Formaldehydeâ€Assisted Isolation of Regulatory Elements (FAIRE). Current Protocols in Molecular Biology, 2013, 102, Unit21.26.	2.9	35
29	Tumor-specific retargeting of an oncogenic transcription factor chimera results in dysregulation of chromatin and transcription. Genome Research, 2012, 22, 259-270.	5.5	96
30	Using formaldehyde-assisted isolation of regulatory elements (FAIRE) to isolate active regulatory DNA. Nature Protocols, 2012, 7, 256-267.	12.0	274
31	MiT Transcription Factor Associated Malignancies in Man. Cell Cycle, 2007, 6, 1724-1729.	2.6	31