

Paul Guilhamon

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1569831/publications.pdf>

Version: 2024-02-01

16
papers

1,465
citations

687363

13
h-index

940533

16
g-index

20
all docs

20
docs citations

20
times ranked

3132
citing authors

#	ARTICLE	IF	CITATIONS
1	Fate mapping of human glioblastoma reveals an invariant stem cell hierarchy. <i>Nature</i> , 2017, 549, 227-232.	27.8	321
2	Integrated (epi)-Genomic Analyses Identify Subgroup-Specific Therapeutic Targets in CNS Rhabdoid Tumors. <i>Cancer Cell</i> , 2016, 30, 891-908.	16.8	191
3	ASCL1 Reorganizes Chromatin to Direct Neuronal Fate and Suppress Tumorigenicity of Glioblastoma Stem Cells. <i>Cell Stem Cell</i> , 2017, 21, 209-224.e7.	11.1	150
4	Gradient of Developmental and Injury Response transcriptional states defines functional vulnerabilities underpinning glioblastoma heterogeneity. <i>Nature Cancer</i> , 2021, 2, 157-173.	13.2	147
5	Pervasive H3K27 Acetylation Leads to ERV Expression and a Therapeutic Vulnerability in H3K27M Gliomas. <i>Cancer Cell</i> , 2019, 35, 782-797.e8.	16.8	143
6	Epigenetic Switchâ€“Induced Viral Mimicry Evasion in Chemotherapy-Resistant Breast Cancer. <i>Cancer Discovery</i> , 2020, 10, 1312-1329.	9.4	84
7	Metabolic Regulation of the Epigenome Drives Lethal Infantile Ependymoma. <i>Cell</i> , 2020, 181, 1329-1345.e24.	28.9	79
8	PRMT5 inhibition disrupts splicing and stemness in glioblastoma. <i>Nature Communications</i> , 2021, 12, 979.	12.8	77
9	A C19MC-LIN28A-MYCN Oncogenic Circuit Driven by Hijacked Super-enhancers Is a Distinct Therapeutic Vulnerability in ETMRs: A Lethal Brain Tumor. <i>Cancer Cell</i> , 2019, 36, 51-67.e7.	16.8	69
10	High-resolution structural genomics reveals new therapeutic vulnerabilities in glioblastoma. <i>Genome Research</i> , 2019, 29, 1211-1222.	5.5	52
11	Cistrome Partitioning Reveals Convergence of Somatic Mutations and Risk Variants on Master Transcription Regulators in Primary Prostate Tumors. <i>Cancer Cell</i> , 2019, 36, 674-689.e6.	16.8	52
12	Single-cell chromatin accessibility profiling of glioblastoma identifies an invasive cancer stem cell population associated with lower survival. <i>ELife</i> , 2021, 10, .	6.0	45
13	Convergence of BMI1 and CHD7 on ERK Signaling in Medulloblastoma. <i>Cell Reports</i> , 2017, 21, 2772-2784.	6.4	31
14	C3D: a tool to predict 3D genomic interactions between cis-regulatory elements. <i>Bioinformatics</i> , 2019, 35, 877-879.	4.1	11
15	SMuRF: a novel tool to identify regulatory elements enriched for somatic point mutations. <i>BMC Bioinformatics</i> , 2018, 19, 454.	2.6	4
16	Mutations in Noncoding <i>Cis</i> -Regulatory Elements Reveal Cancer Driver Cistromes in Luminal Breast Cancer. <i>Molecular Cancer Research</i> , 2022, 20, 102-113.	3.4	3