

Luciano Garofano

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

5,365
citations

932766

10
h-index

1125271

13
g-index

20
all docs

20
docs citations

20
times ranked

9765
citing authors

#	ARTICLE	IF	CITATIONS
1	Regulated interaction of ID2 with the anaphase-promoting complex links progression through mitosis with reactivation of cell-type-specific transcription. <i>Nature Communications</i> , 2022, 13, 2089.	5.8	2
2	Pathway-based classification of glioblastoma uncovers a mitochondrial subtype with therapeutic vulnerabilities. <i>Nature Cancer</i> , 2021, 2, 141-156.	5.7	163
3	Proteogenomic insights into the biology and treatment of HPV-negative head and neck squamous cell carcinoma. <i>Cancer Cell</i> , 2021, 39, 361-379.e16.	7.7	189
4	Proteogenomic and metabolomic characterization of human glioblastoma. <i>Cancer Cell</i> , 2021, 39, 509-528.e20.	7.7	327
5	OMRT-3. Longitudinal analysis of diffuse glioma reveals cell state dynamics at recurrence associated with changes in genetics and the microenvironment. <i>Neuro-Oncology Advances</i> , 2021, 3, ii7-ii8.	0.4	1
6	OTEH-10. Evolutionary trajectory of epigenomic of gliomas. <i>Neuro-Oncology Advances</i> , 2021, 3, ii12-ii12.	0.4	0
7	EPCO-09. LONGITUDINAL ANALYSIS OF DIFFUSE GLIOMA REVEALS CELL STATE DYNAMICS AT RECURRENCE ASSOCIATED WITH CHANGES IN GENETICS AND THE MICROENVIRONMENT. <i>Neuro-Oncology</i> , 2021, 23, vi3-vi3.	0.6	0
8	TAMI-52. NEURONAL MECHANISMS OF BRAIN TUMOR INVASION. <i>Neuro-Oncology</i> , 2021, 23, vi209-vi209.	0.6	0
9	Temporospatial genomic profiling in glioblastoma identifies commonly altered core pathways underlying tumor progression. <i>Neuro-Oncology Advances</i> , 2020, 2, vdaa078.	0.4	12
10	A map of tumor-host interactions in glioma at single-cell resolution. <i>GigaScience</i> , 2020, 9, .	3.3	32
11	Treatment of metabolic acidosis with sodium bicarbonate delays progression of chronic kidney disease: the UBI Study. <i>Journal of Nephrology</i> , 2019, 32, 989-1001.	0.9	104
12	The molecular landscape of glioma in patients with Neurofibromatosis 1. <i>Nature Medicine</i> , 2019, 25, 176-187.	15.2	145
13	RGBM: regularized gradient boosting machines for identification of the transcriptional regulators of discrete glioma subtypes. <i>Nucleic Acids Research</i> , 2018, 46, e39-e39.	6.5	32
14	A metabolic function of FGFR3-TACC3 gene fusions in cancer. <i>Nature</i> , 2018, 553, 222-227.	13.7	137
15	Molecular Profiling Reveals Biologically Discrete Subsets and Pathways of Progression in Diffuse Glioma. <i>Cell</i> , 2016, 164, 550-563.	13.5	1,695
16	TCGAbiolinks: an R/Bioconductor package for integrative analysis of TCGA data. <i>Nucleic Acids Research</i> , 2016, 44, e71-e71.	6.5	2,519