

Giacomo Corleone

List of Publications by Year in descending order

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14
papers

1,380
citations

933447

10
h-index

1125743

13
g-index

19
all docs

19
docs citations

19
times ranked

2520
citing authors

#	ARTICLE	IF	CITATIONS
1	KEAP1 and TP53 Frame Genomic, Evolutionary, and Immunologic Subtypes of Lung Adenocarcinoma With Different Sensitivity to Immunotherapy. <i>Journal of Thoracic Oncology</i> , 2021, 16, 2065-2077.	1.1	28
2	KEAP1-driven co-mutations in lung adenocarcinoma unresponsive to immunotherapy despite high tumor mutational burden. <i>Annals of Oncology</i> , 2020, 31, 1746-1754.	1.2	140
3	Che-1/AATF-induced transcriptionally active chromatin promotes cell proliferation in multiple myeloma. <i>Blood Advances</i> , 2020, 4, 5616-5630.	5.2	10
4	GeDi: applying suffix arrays to increase the repertoire of detectable SNVs in tumour genomes. <i>BMC Bioinformatics</i> , 2020, 21, 45.	2.6	0
5	Eleven grand challenges in single-cell data science. <i>Genome Biology</i> , 2020, 21, 31.	8.8	742
6	Small extracellular vesicles deliver miRâ€21 and miRâ€17 as proâ€senescence effectors to endothelial cells. <i>Journal of Extracellular Vesicles</i> , 2020, 9, 1725285.	12.2	104
7	B4GALT1 Is a New Candidate to Maintain the Stemness of Lung Cancer Stem Cells. <i>Journal of Clinical Medicine</i> , 2019, 8, 1928.	2.4	13
8	Single-cell transcriptomics reveals multi-step adaptations to endocrine therapy. <i>Nature Communications</i> , 2019, 10, 3840.	12.8	93
9	Adhesion-Mediated Multiple Myeloma (MM) Disease Progression: Junctional Adhesion Molecule a Enhances Angiogenesis and Multiple Myeloma Dissemination and Predicts Poor Survival. <i>Blood</i> , 2019, 134, 855-855.	1.4	7
10	Extensive and systematic rewiring of histone post-translational modifications in cancer model systems. <i>Nucleic Acids Research</i> , 2018, 46, 3817-3832.	14.5	31
11	Chromatin Immunoprecipitation and High-Throughput Sequencing (ChIP-Seq): Tips and Tricks Regarding the Laboratory Protocol and Initial Downstream Data Analysis. <i>Methods in Molecular Biology</i> , 2018, 1767, 271-288.	0.9	2
12	Enhancer mapping uncovers phenotypic heterogeneity and evolution in patients with luminal breast cancer. <i>Nature Medicine</i> , 2018, 24, 1469-1480.	30.7	98
13	Acquired CYP19A1 amplification is an early specific mechanism of aromatase inhibitor resistance in ERâ€ metastatic breast cancer. <i>Nature Genetics</i> , 2017, 49, 444-450.	21.4	77
14	Large-scale profiling of signalling pathways reveals an asthma specific signature in bronchial smooth muscle cells. <i>Oncotarget</i> , 2016, 7, 25150-25161.	1.8	32