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List of Publications by Year in descending order

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Version: 2024-02-01

933447 1125743 1,380 14 10 13 citations g-index h-index papers 19 19 19 2520 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Eleven grand challenges in single-cell data science. Genome Biology, 2020, 21, 31.	8.8	742
2	KEAP1-driven co-mutations in lung adenocarcinoma unresponsive to immunotherapy despite high tumor mutational burden. Annals of Oncology, 2020, 31, 1746-1754.	1.2	140
3	Small extracellular vesicles deliver miRâ€21 and miRâ€217 as proâ€senescence effectors to endothelial cells. Journal of Extracellular Vesicles, 2020, 9, 1725285.	12.2	104
4	Enhancer mapping uncovers phenotypic heterogeneity and evolution in patients with luminal breast cancer. Nature Medicine, 2018, 24, 1469-1480.	30.7	98
5	Single-cell transcriptomics reveals multi-step adaptations to endocrine therapy. Nature Communications, 2019, 10, 3840.	12.8	93
6	Acquired CYP19A1 amplification is an early specific mechanism of aromatase inhibitor resistance in ER $\hat{l}\pm$ metastatic breast cancer. Nature Genetics, 2017, 49, 444-450.	21.4	77
7	Large-scale profiling of signalling pathways reveals an asthma specific signature in bronchial smooth muscle cells. Oncotarget, 2016, 7, 25150-25161.	1.8	32
8	Extensive and systematic rewiring of histone post-translational modifications in cancer model systems. Nucleic Acids Research, 2018, 46, 3817-3832.	14.5	31
9	KEAP1 and TP53 Frame Genomic, Evolutionary, and Immunologic Subtypes of Lung Adenocarcinoma With Different Sensitivity to Immunotherapy. Journal of Thoracic Oncology, 2021, 16, 2065-2077.	1.1	28
10	B4GALT1 Is a New Candidate to Maintain the Stemness of Lung Cancer Stem Cells. Journal of Clinical Medicine, 2019, 8, 1928.	2.4	13
11	Che-1/AATF-induced transcriptionally active chromatin promotes cell proliferation in multiple myeloma. Blood Advances, 2020, 4, 5616-5630.	5.2	10
12	Adhesion-Mediated Multiple Myeloma (MM) Disease Progression: Junctional Adhesion Molecule a Enhances Angiogenesis and Multiple Myeloma Dissemination and Predicts Poor Survival. Blood, 2019, 134, 855-855.	1.4	7
13	Chromatin Immunoprecipitation and High-Throughput Sequencing (ChIP-Seq): Tips and Tricks Regarding the Laboratory Protocol and Initial Downstream Data Analysis. Methods in Molecular Biology, 2018, 1767, 271-288.	0.9	2
14	GeDi: applying suffix arrays to increase the repertoire of detectable SNVs in tumour genomes. BMC Bioinformatics, 2020, 21, 45.	2.6	0