

Yueli Cui

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3780586/publications.pdf>

Version: 2024-02-01

9
papers

1,127
citations

1162889

8
h-index

1474057

9
g-index

9
all docs

9
docs citations

9
times ranked

2233
citing authors

#	ARTICLE	IF	CITATIONS
1	Heterogeneity of glial progenitor cells during the neurogenesis-to-gliogenesis switch in the developing human cerebral cortex. <i>Cell Reports</i> , 2021, 34, 108788.	2.9	55
2	Single-cell transcriptome and genome analyses of pituitary neuroendocrine tumors. <i>Neuro-Oncology</i> , 2021, 23, 1859-1871.	0.6	29
3	Single-cell analysis of ploidy and the transcriptome reveals functional and spatial divergency in murine megakaryopoiesis. <i>Blood</i> , 2021, 138, 1211-1224.	0.6	59
4	Single-cell transcriptomics identifies divergent developmental lineage trajectories during human pituitary development. <i>Nature Communications</i> , 2020, 11, 5275.	5.8	79
5	Single-Cell Multiomics Sequencing Reveals Prevalent Genomic Alterations in Tumor Stromal Cells of Human Colorectal Cancer. <i>Cancer Cell</i> , 2020, 38, 818-828.e5.	7.7	146
6	Single-Cell Transcriptome Analysis Maps the Developmental Track of the Human Heart. <i>Cell Reports</i> , 2019, 26, 1934-1950.e5.	2.9	355
7	Transcriptional and Spatial Heterogeneity of Mouse Megakaryocytes at Single-Cell Resolution. <i>Blood</i> , 2019, 134, 275-275.	0.6	4
8	Dissecting the Global Dynamic Molecular Profiles of Human Fetal Kidney Development by Single-Cell RNA Sequencing. <i>Cell Reports</i> , 2018, 24, 3554-3567.e3.	2.9	91
9	Single-Cell RNA Sequencing Analysis Reveals Sequential Cell Fate Transition during Human Spermatogenesis. <i>Cell Stem Cell</i> , 2018, 23, 599-614.e4.	5.2	309