Kaikun Xie

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/370948/kaikun-xie-publications-by-year.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

4	69	1	5
papers	citations	h-index	g-index
5	123	8.3 avg, IF	1.46
ext. papers	ext. citations		L-index

#	Paper Control of the	IF	Citations
4	redPATH: Reconstructing the Pseudo Development Time of Cell Lineages in Single-cell RNA-seq Data and Applications in Cancer. <i>Genomics, Proteomics and Bioinformatics</i> , 2021 , 19, 292-305	6.5	O
3	scAIDE: clustering of large-scale single-cell RNA-seq data reveals putative and rare cell types. <i>NAR Genomics and Bioinformatics</i> , 2020 , 2, lqaa082	3.7	1
2	Reconstructing cell cycle pseudo time-series via single-cell transcriptome data. <i>Nature Communications</i> , 2017 , 8, 22	17.4	67
1	redPATH: Reconstructing the Pseudo Development Time of Cell Lineages in Single-Cell RNA-Seq Data and Applications in Cancer		1