## Kaikun Xie

## List of Publications by Citations

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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 g-index

5 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
4	Reconstructing cell cycle pseudo time-series via single-cell transcriptome data. <i>Nature Communications</i> , <b>2017</b> , 8, 22	17.4	67
3	scAIDE: clustering of large-scale single-cell RNA-seq data reveals putative and rare cell types. <i>NAR Genomics and Bioinformatics</i> , <b>2020</b> , 2, lqaa082	3.7	1
2	redPATH: Reconstructing the Pseudo Development Time of Cell Lineages in Single-Cell RNA-Seq Data and Applications in Cancer		1
1	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> <b>2021</b> , 19, 292-305	6.5	0