## Tianyu Zhu

## 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.

6 83 5 9 g-index

9 151 9.4 2.51 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
6	Cell and tissue type independent age-associated DNA methylation changes are not rare but common. <i>Aging</i> , <b>2018</b> , 10, 3541-3557	5.6	28
5	EPISCORE: cell type deconvolution of bulk tissue DNA methylomes from single-cell RNA-Seq data. <i>Genome Biology</i> , <b>2020</b> , 21, 221	18.3	21
4	A cell-type deconvolution meta-analysis of whole blood EWAS reveals lineage-specific smoking-associated DNA methylation changes. <i>Nature Communications</i> , <b>2020</b> , 11, 4779	17.4	14
3	EpiDISH web server: Epigenetic Dissection of Intra-Sample-Heterogeneity with online GUI. <i>Bioinformatics</i> , <b>2019</b> ,	7.2	10
2	Clinical outcomes, Kadish-INSICA staging and therapeutic targeting of somatostatin receptor 2 in olfactory neuroblastoma <i>European Journal of Cancer</i> , <b>2021</b> ,	7.5	5
1	A pan-tissue DNA methylation atlas enables in silico decomposition of human tissue methylomes at cell-type resolution <i>Nature Methods</i> , <b>2022</b> , 19, 296-306	21.6	3