Qi Tian

List of Publications by Year
in descending order

8 papers	**67** citations	2258059 **3** h-index	1720034 **7** g-index
8 all docs	**8** docs citations	**8** times ranked	**63** citing authors

#	Article	IF	Citations
1	MRCNN: a deep learning model for regression of genome-wide DNA methylation. BMC Genomics, 2019, 20, 192.	2.8	36
2	CaMelia: imputation in single-cell methylomes based on local similarities between cells. Bioinformatics, 2021, 37, 1814-1820.	4.1	12
3	A robust fuzzy rule based integrative feature selection strategy for gene expression data in TCGA. BMC Medical Genomics, 2019, 12, 14.	1.5	5
4	PretiMeth: precise prediction models for DNA methylation based on single methylation mark. BMC Genomics, 2020, 21, 364.	2.8	5
5	A Hybrid Ensemble Approach for Identifying Robust Differentially Methylated Loci in Pan-Cancers. Frontiers in Genetics, 2019, 10, 774.	2.3	4
6	scMelody: An Enhanced Consensus-Based Clustering Model for Single-Cell Methylation Data by Reconstructing Cell-to-Cell Similarity. Frontiers in Bioengineering and Biotechnology, 2022, 10, 842019.	4.1	4
7	The progress on the estimation of DNA methylation level and the detection of abnormal methylation. Quantitative Biology, 2022, 10, 55-66.	0.5	1
8	Imputation in Scrna-seq Data Using Supervised Deep Generative Networks. , 2021, , .		0