

# Qi Tian

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9848957/publications.pdf>

Version: 2024-02-01

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