

# Hamim Zafar

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3169793/publications.pdf>

Version: 2024-02-01

7  
papers

469  
citations

1684188

5  
h-index

1872680

6  
g-index

10  
all docs

10  
docs citations

10  
times ranked

631  
citing authors

#	ARTICLE	IF	CITATIONS
1	SiFit: inferring tumor trees from single-cell sequencing data under finite-sites models. <i>Genome Biology</i> , 2017, 18, 178.	8.8	152
2	Monovar: single-nucleotide variant detection in single cells. <i>Nature Methods</i> , 2016, 13, 505-507.	19.0	150
3	SiCloneFit: Bayesian inference of population structure, genotype, and phylogeny of tumor clones from single-cell genome sequencing data. <i>Genome Research</i> , 2019, 29, 1847-1859.	5.5	97
4	Computational approaches for inferring tumor evolution from single-cell genomic data. <i>Current Opinion in Systems Biology</i> , 2018, 7, 16-25.	2.6	36
5	Matrilysin/MMP-7 Cleavage of Perlecan/HSPG2 Complexed with Semaphorin 3A Supports FAK-Mediated Stromal Invasion by Prostate Cancer Cells. <i>Scientific Reports</i> , 2018, 8, 7262.	3.3	25
6	Inference of cell state transitions and cell fate plasticity from single-cell with MARGARET. <i>Nucleic Acids Research</i> , 2022, 50, e86-e86.	14.5	6
7	Comments on the model parameters in "SiFit: inferring tumor trees from single-cell sequencing data under finite-sites models", <i>Genome Biology</i> , 2019, 20, 95.	8.8	1