

# Shu-Lin Wang

## List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Are dropout imputation methods for scRNA-seq effective for scATAC-seq data?. Briefings in Bioinformatics, 2022, 23, .	6.5	5
2	DMFMDA: Prediction of Microbe-Disease Associations Based on Deep Matrix Factorization Using Bayesian Personalized Ranking. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2021, 18, 1763-1772.	3.0	21
3	An Integrated Framework for Identifying Mutated Driver Pathway and Cancer Progression. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2019, 16, 455-464.	3.0	12
4	Prediction of Microbe-Disease Associations by Graph Regularized Non-Negative Matrix Factorization. Journal of Computational Biology, 2018, 25, 1385-1394.	1.6	16