

# Menglan Cai

## List of Publications by Year in descending order

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citing authors

#	ARTICLE	IF	CITATIONS
1	XGSEA: CROSS-species gene set enrichment analysis via domain adaptation. Briefings in Bioinformatics, 2021, 22, .	6.5	9
2	MicroRNA-disease association prediction by matrix tri-factorization. BMC Genomics, 2020, 21, 617.	2.8	4
3	Cross-species Data Classification by Domain Adaptation via Discriminative Heterogeneous Maximum Mean Discrepancy. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2019, 18, 1-1.	3.0	7
4	Drug repositioning via matrix completion with multi-view side information. IET Systems Biology, 2019, 13, 267-275.	1.5	2
5	Integrative subspace clustering by common and specific decomposition for applications on cancer subtype identification. BMC Medical Genomics, 2019, 12, 191.	1.5	4
6	Drug Target Prediction by Multi-View Low Rank Embedding. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2019, 16, 1712-1721.	3.0	17
7	rPCMP: robust p-value combination by multiple partitions with applications to ATAC-seq data. BMC Systems Biology, 2018, 12, 141.	3.0	0
8	Subtype identification from heterogeneous TCGA datasets on a genomic scale by multi-view clustering with enhanced consensus. BMC Medical Genomics, 2017, 10, 75.	1.5	16