Mi-Xiao Hou

List of Publications by Year
in descending order

14 papers	92 citations	1684188 5 h-index	1372567 10 g-index
14 all docs	14 docs citations	14 times ranked	73 citing authors

#	Article	IF	Citations
1	Multi-View Speech Emotion Recognition Via Collective Relation Construction. IEEE/ACM Transactions on Audio Speech and Language Processing, 2022, 30, 218-229.	5.8	23
2	Multimodal Emotion Recognition With Temporal and Semantic Consistency. IEEE/ACM Transactions on Audio Speech and Language Processing, 2021, 29, 3592-3603.	5.8	18
3	Robust Nonnegative Matrix Factorization via Joint Graph Laplacian and Discriminative Information for Identifying Differentially Expressed Genes. Complexity, 2017, 2017, 1-11.	1.6	16
4	An Integrated Graph Regularized Non-Negative Matrix Factorization Model for Gene Co-Expression Network Analysis. IEEE Access, 2019, 7, 126594-126602.	4.2	9
5	A new method for mining information of co-expression network based on multi-cancers integrated data. BMC Medical Genomics, 2019, 12, 155.	1.5	6
6	Network analysis based on low-rank method for mining information on integrated data of multi-cancers. Computational Biology and Chemistry, 2019, 78, 468-473.	2.3	5
7	Multi-Modal Emotion Recognition with Self-Guided Modality Calibration. , 2022, , .		5
8	Hypergraph Regularized Discriminative Nonnegative Matrix Factorization on Sample Classification and Co-Differentially Expressed Gene Selection. Complexity, 2019, 2019, 1-12.	1.6	4
9	Application of a Deep Matrix Factorization Model on Integrated Gene Expression Data. Current Bioinformatics, 2020, 15, 359-367.	1.5	4
10	LJELSR: A Strengthened Version of JELSR for Feature Selection and Clustering. International Journal of Molecular Sciences, 2019, 20, 886.	4.1	1
11	A New Model of Identifying Differentially Expressed Genes via Weighted Network Analysis Based on Dimensionality Reduction Method. Current Bioinformatics, 2019, 14, 762-770.	1.5	1
12	Robust graph regularized discriminative nonnegative matrix factorization for characteristic gene selection. , 2016, , .		0
13	A p-norm singular value decomposition method for robust tumor clustering. , 2016, , .		0
14	Sparse Orthogonal Nonnegative Matrix Factorization for Identifying Differentially Expressed Genes and Clustering Tumor Samples. , 2018, , .		0