

# Mi-Xiao Hou

## List of Publications by Year in descending order

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Version: 2024-02-01

14  
papers

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citations

1684188

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1372567

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docs citations

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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	Multi-Modal Emotion Recognition with Self-Guided Modality Calibration. , 2022, , .		5
3	Multimodal Emotion Recognition With Temporal and Semantic Consistency. IEEE/ACM Transactions on Audio Speech and Language Processing, 2021, 29, 3592-3603.	5.8	18
4	Application of a Deep Matrix Factorization Model on Integrated Gene Expression Data. Current Bioinformatics, 2020, 15, 359-367.	1.5	4
5	An Integrated Graph Regularized Non-Negative Matrix Factorization Model for Gene Co-Expression Network Analysis. IEEE Access, 2019, 7, 126594-126602.	4.2	9
6	Hypergraph Regularized Discriminative Nonnegative Matrix Factorization on Sample Classification and Co-Differentially Expressed Gene Selection. Complexity, 2019, 2019, 1-12.	1.6	4
7	LJELSR: A Strengthened Version of JELSR for Feature Selection and Clustering. International Journal of Molecular Sciences, 2019, 20, 886.	4.1	1
8	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
9	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
10	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
11	Sparse Orthogonal Nonnegative Matrix Factorization for Identifying Differentially Expressed Genes and Clustering Tumor Samples. , 2018, , .		0
12	Robust Nonnegative Matrix Factorization via Joint Graph Laplacian and Discriminative Information for Identifying Differentially Expressed Genes. Complexity, 2017, 2017, 1-11.	1.6	16
13	Robust graph regularized discriminative nonnegative matrix factorization for characteristic gene selection. , 2016, , .		0
14	A p-norm singular value decomposition method for robust tumor clustering. , 2016, , .		0