

Na Yu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2947564/publications.pdf>

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10
papers

170
citations

1478505

6
h-index

1720034

7
g-index

11
all docs

11
docs citations

11
times ranked

120
citing authors

#	ARTICLE	IF	CITATIONS
1	Predicting multiple types of MicroRNA-disease associations based on tensor factorization and label propagation. <i>Computers in Biology and Medicine</i> , 2022, 146, 105558.	7.0	8
2	Correntropy-Based Hypergraph Regularized NMF for Clustering and Feature Selection on Multi-Cancer Integrated Data. <i>IEEE Transactions on Cybernetics</i> , 2021, 51, 3952-3963.	9.5	62
3	Dual Hyper-Graph Regularized Supervised NMF for Selecting Differentially Expressed Genes and Tumor Classification. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2021, 18, 2375-2383.	3.0	12
4	GSEMT: A Gene Set Enrichment Analysis Method Based on Mantel Test. <i>Journal of Physics: Conference Series</i> , 2021, 1828, 012048.	0.4	0
5	Hyper-Graph Regularized Constrained NMF for Selecting Differentially Expressed Genes and Tumor Classification. <i>IEEE Journal of Biomedical and Health Informatics</i> , 2020, 24, 3002-3011.	6.3	37
6	Robust hypergraph regularized non-negative matrix factorization for sample clustering and feature selection in multi-view gene expression data. <i>Human Genomics</i> , 2019, 13, 46.	2.9	8
7	Sparse Graph Regularization Non-Negative Matrix Factorization Based on Huber Loss Model for Cancer Data Analysis. <i>Frontiers in Genetics</i> , 2019, 10, 1054.	2.3	7
8	Hypergraph regularized NMF by $L_{2,1}$ -norm for Clustering and Com-abnormal Expression Genes Selection. , 2018, , .		4
9	Co-differential Gene Selection and Clustering Based on Graph Regularized Multi-View NMF in Cancer Genomic Data. <i>Genes</i> , 2018, 9, 586.	2.4	25
10	Graph regularized robust non-negative matrix factorization for clustering and selecting differentially expressed genes. , 2017, , .		5