

Li Huang

List of Publications by Year in descending order

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11
papers

1,308
citations

1040056

9
h-index

1281871

11
g-index

11
all docs

11
docs citations

11
times ranked

971
citing authors

#	ARTICLE	IF	CITATIONS
1	Prediction of potential miRNA-disease associations based on stacked autoencoder. Briefings in Bioinformatics, 2022, 23, .	6.5	31
2	Identification of miRNA-disease associations via deep forest ensemble learning based on autoencoder. Briefings in Bioinformatics, 2022, 23, .	6.5	60
3	RNMFLP: Predicting circRNA-disease associations based on robust nonnegative matrix factorization and label propagation. Briefings in Bioinformatics, 2022, 23, .	6.5	32
4	Fusion of KATZ measure and space projection to fast probe potential lncRNA-disease associations in bipartite graphs. PLoS ONE, 2021, 16, e0260329.	2.5	1
5	In silico prediction of potential miRNA-disease association using an integrative bioinformatics approach based on kernel fusion. Journal of Cellular and Molecular Medicine, 2020, 24, 573-587.	3.6	8
6	LoAdaBoost: Loss-based AdaBoost federated machine learning with reduced computational complexity on IID and non-IID intensive care data. PLoS ONE, 2020, 15, e0230706.	2.5	80
7	Patient clustering improves efficiency of federated machine learning to predict mortality and hospital stay time using distributed electronic medical records. Journal of Biomedical Informatics, 2019, 99, 103291.	4.3	229
8	EGBMMDA: Extreme Gradient Boosting Machine for MiRNA-Disease Association prediction. Cell Death and Disease, 2018, 9, 3.	6.3	256
9	NDAMDA: Network distance analysis for MiRNA-disease association prediction. Journal of Cellular and Molecular Medicine, 2018, 22, 2884-2895.	3.6	34
10	MDHGI: Matrix Decomposition and Heterogeneous Graph Inference for miRNA-disease association prediction. PLoS Computational Biology, 2018, 14, e1006418.	3.2	323
11	LRSSLMDA: Laplacian Regularized Sparse Subspace Learning for MiRNA-Disease Association prediction. PLoS Computational Biology, 2017, 13, e1005912.	3.2	254