Minhyeok Lee

List of Publications by Year
in descending order

14 papers	140 citations	1478280 6 h-index	1281743 11 g-index
14 all docs	14 docs citations	14 times ranked	95 citing authors

#	Article	IF	Citations
1	HRGAN: A Generative Adversarial Network Producing Higher-Resolution Images than Training Sets. Sensors, 2022, 22, 1435.	2.1	7
2	An Ensemble Deep Learning Model with a Gene Attention Mechanism for Estimating the Prognosis of Low-Grade Glioma. Biology, 2022, 11, 586.	1.3	7
3	Estimation with Uncertainty via Conditional Generative Adversarial Networks. Sensors, 2021, 21, 6194.	2.1	9
4	Portfolio management via two-stage deep learning with a joint cost. Expert Systems With Applications, 2020, 143, 113041.	4.4	27
5	Long Term Traffic Prediction in Highway Using Parallel CNN. , 2020, , .		2
6	Improved recurrent generative adversarial networks with regularization techniques and a controllable framework. Information Sciences, 2020, 538, 428-443.	4.0	7
7	Serum biomarkers from cell-based assays for AhRL and MIS strongly predicted the future development of diabetes in a large community-based prospective study in Korea. Scientific Reports, 2020, 10, 6339.	1.6	9
8	Clinical Implication of Concordant or Discordant Genomic Profiling between Primary and Matched Metastatic Tissues in Patients with Colorectal Cancer. Cancer Research and Treatment, 2020, 52, 764-778.	1.3	6
9	Stock Price Prediction Through the Sentimental Analysis of News Articles. , 2019, , .		15
10	Controllable Generative Adversarial Network. IEEE Access, 2019, 7, 28158-28169.	2.6	45
11	Prediction of survival risks with adjusted gene expression through risk-gene networks. Bioinformatics, 2019, 35, 4898-4906.	1.8	1
12	Regression-Based Network Estimation for High-Dimensional Genetic Data. Journal of Computational Biology, 2019, 26, 336-349.	0.8	1
13	The estimation of probability distribution for factor variables with many categorical values. PLoS ONE, 2018, 13, e0202547.	1.1	2
14	A Comparison of Two-Stage Approaches Based on Penalized Regression for Estimating Gene Networks. Journal of Computational Biology, 2017, 24, 709-720.	0.8	2