

Jnanendra Prasad Sarkar

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

232
citations

1478505

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h-index

1125743

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g-index

19
all docs

19
docs citations

19
times ranked

325
citing authors

#	ARTICLE	IF	CITATIONS
1	Human miRNAs to Identify Potential Regions of SARS-CoV-2. ACS Omega, 2022, 7, 21086-21101.	3.5	2
2	Online Predictor Using Machine Learning to Predict Novel Coronavirus and Other Pathogenic Viruses. ACS Omega, 2022, 7, 23069-23074.	3.5	0
3	Topological Analysis for Sequence Variability: Case Study on more than 2K SARS-CoV-2 sequences of COVID-19 infected 54 countries in comparison with SARS-CoV-1 and MERS-CoV. Infection, Genetics and Evolution, 2021, 88, 104708.	2.3	6
4	Machine learning integrated ensemble of feature selection methods followed by survival analysis for predicting breast cancer subtype specific miRNA biomarkers. Computers in Biology and Medicine, 2021, 131, 104244.	7.0	49
5	Strategies for COVID-19 Epidemiological Surveillance in India: Overall Policies Till June 2021. Frontiers in Public Health, 2021, 9, 708224.	2.7	2
6	Machine learning integrated credibilistic semi supervised clustering for categorical data. Applied Soft Computing Journal, 2020, 86, 105871.	7.2	5
7	Genome-wide analysis of Indian SARS-CoV-2 genomes for the identification of genetic mutation and SNP. Infection, Genetics and Evolution, 2020, 85, 104457.	2.3	35
8	Identification of miRNA Biomarkers for Diverse Cancer Types Using Statistical Learning Methods at the Whole-Genome Scale. Frontiers in Genetics, 2020, 11, 982.	2.3	4
9	A new evolutionary rough fuzzy integrated machine learning technique for microRNA selection using next-generation sequencing data of breast cancer. , 2019, , .		1
10	Improved Fuzzy Clustering using Ensemble based Differential Evolution for Remote Sensing Image. , 2019, , .		1
11	Identification of Breast Cancer Subtype Specific MicroRNAs Using Survival Analysis to Find Their Role in Transcriptomic Regulation. Frontiers in Genetics, 2019, 10, 1047.	2.3	16
12	Integrated Rough Fuzzy Clustering for Categorical data Analysis. Fuzzy Sets and Systems, 2019, 361, 1-32.	2.7	26
13	Survival Analysis with the Integration of RNA-Seq and Clinical Data to Identify Breast Cancer Subtype Specific Genes. Lecture Notes in Computer Science, 2019, , 139-146.	1.3	1
14	Improving Modified Differential Evolution for Fuzzy Clustering. Advances in Intelligent Systems and Computing, 2018, , 136-146.	0.6	1
15	Rough Possibilistic Type-2 Fuzzy C-Means clustering for MR brain image segmentation. Applied Soft Computing Journal, 2016, 46, 527-536.	7.2	53
16	Ensemble based rough fuzzy clustering for categorical data. Knowledge-Based Systems, 2015, 77, 114-127.	7.1	28
17	A new SVM integrated rough type-II fuzzy clustering technique. , 2014, , .		0
18	Rough Set Based Fuzzy K-Modes for Categorical Data. Lecture Notes in Computer Science, 2012, , 323-330.	1.3	2