

Zahra Mungloo-Dilmohamud

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

125
citations

1684188

5
h-index

1372567

10
g-index

24
all docs

24
docs citations

24
times ranked

227
citing authors

#	ARTICLE	IF	CITATIONS
1	VAMDLE: Visitor and Asset Management Using Deep Learning and ElasticSearch. Lecture Notes in Networks and Systems, 2022, , 318-329.	0.7	0
2	Balancing Data through Data Augmentation Improves the Generality of Transfer Learning for Diabetic Retinopathy Classification. Applied Sciences (Switzerland), 2022, 12, 5363.	2.5	8
3	A Deep Transfer Learning Model for the Identification of Bird Songs: A Case Study for Mauritius. , 2021, , .		0
4	Ensemble Feature Selection: Are Stability Metrics a Proxy or a Complement to Predictive Performance?. , 2021, , .		0
5	Multi-Objective Optimisation for Improved Feature Selection. , 2021, , .		0
6	Recent advances in computational tools and resources for the self-management of type 2 diabetes. Informatics for Health and Social Care, 2020, 45, 77-95.	2.6	8
7	DOT: a crowdsourcing Mobile application for disease outbreak detection and surveillance in Mauritius. Health and Technology, 2020, 10, 1115-1127.	3.6	0
8	Stability of Feature Selection Methods: A Study of Metrics Across Different Gene Expression Datasets. Lecture Notes in Computer Science, 2020, , 659-669.	1.3	2
9	Job Recommendation System, Machine Learning, Regression, Classification, Natural Language Processing. , 2020, , .		6
10	Health Data Analytics: Current Perspectives, Challenges, and Future Directions. EAI/Springer Innovations in Communication and Computing, 2020, , 117-151.	1.1	0
11	Discovering novel driver mutations from pan-cancer analysis of mutational and gene expression profiles. PLoS ONE, 2020, 15, e0242780.	2.5	6
12	EyeSee: Camera to Caption with Attention Mechanism. , 2020, , .		0
13	Detection of Genome Sequence Outliers Across Pan-Genomes. , 2019, , .		0
14	Proposed minimum information guideline for kidney disease research and clinical data reporting: a cross-sectional study. BMJ Open, 2019, 9, e029539.	1.9	1
15	Exploring the Stability of Feature Selection Methods across a Palette of Gene Expression Datasets. , 2019, , .		4
16	Comparison of genomic islands in cyanobacteria: Evidence of bacteriophage-mediated horizontal gene transfer from eukaryotes. Microbiological Research, 2018, 211, 31-46.	5.3	11
17	A Comparative Study of Feature Selection Methods for Biomarker Discovery. , 2018, , .		3
18	A Meta-Review of Feature Selection Techniques in the Context of Microarray Data. Lecture Notes in Computer Science, 2017, , 33-49.	1.3	5

#	ARTICLE	IF	CITATIONS
19	Development of Bioinformatics Infrastructure for Genomics Research. Global Heart, 2017, 12, 91.	2.3	47
20	Genomic Research Data Generation, Analysis and Sharing – Challenges in the African Setting. Data Science Journal, 2017, 16, .	1.3	21
21	TAAU: A tool to study the amino acid usage in different organisms. , 2016, , .		0
22	H3ABioNet computational metagenomics workshop in Mauritius: training to analyse microbial diversity for Africa. Standards in Genomic Sciences, 2015, 10, .	1.5	2