

Wenjuan Cui

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3285527/publications.pdf>

Version: 2024-02-01

15
papers

91
citations

1478505

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1588992

8
g-index

17
all docs

17
docs citations

17
times ranked

131
citing authors

#	ARTICLE	IF	CITATIONS
1	Machine Learning Prediction of Foodborne Disease Pathogens: Algorithm Development and Validation Study. JMIR Medical Informatics, 2021, 9, e24924.	2.6	12
2	High-Efficiency Machine Learning Method for Identifying Foodborne Disease Outbreaks and Confounding Factors. Foodborne Pathogens and Disease, 2021, 18, 590-598.	1.8	8
3	Foodborne Disease Risk Prediction Using Multigraph Structural Long Short-term Memory Networks: Algorithm Design and Validation Study. JMIR Medical Informatics, 2021, 9, e29433.	2.6	6
4	Spatiotemporal Trend Analysis of PM2.5 Concentration in China, 1999–2016. Atmosphere, 2019, 10, 461.	2.3	7
5	Power optimization through peripheral circuit reusing integrated with loop tiling for RRAM crossbar-based CNN. , 2018, , .		2
6	The Utility of Supertype Clustering in Prediction for Class II MHC-Peptide Binding. Molecules, 2018, 23, 3034.	3.8	9
7	RPiRLS: Quantitative Predictions of RNA Interacting with Any Protein of Known Sequence. Molecules, 2018, 23, 540.	3.8	10
8	A Polynomial Time Approximation Scheme for the Closest Shared Center Problem. Algorithmica, 2017, 77, 65-83.	1.3	0
9	An algorithm for event detection based on social media data. Neurocomputing, 2017, 254, 53-58.	5.9	27
10	Personalized microblog recommendation using sentimental features. , 2017, , .		2
11	Distributed retrieval for massive remote sensing image metadata on spark. , 2016, , .		5
12	How to use open source data to assess infection disease risk: A framework and applications. , 2015, , .		0
13	Self-adaptive Wi-Fi indoor positioning model. , 2015, , .		0
14	A Polynomial Time Approximation Scheme for the Closest Shared Center Problem. Lecture Notes in Computer Science, 2013, , 385-396.	1.3	0
15	Identifying mutation regions for closely related individuals without a known pedigree. BMC Bioinformatics, 2012, 13, 146.	2.6	3